

SUPPLEMENTARY MATERIAL

Isolation and structural elucidation of new amphidinol analogues from *Amphidinium carterae* cultivated in a pilot scale photobioreactor

Adrián Morales-Amador^{1,2}, Alejandro Molina-Miras^{3,4}, Lorenzo López-Rosales^{3,4}, Asterio Sánchez-Mirón^{3,4}, Francisco García-Camacho^{3,4}, María L. Souto^{1,2*} and José J. Fernández^{1,2*}

¹ Instituto Universitario de Bio-Organica Antonio González (IUBO AG), Universidad de La Laguna (ULL), Avda. Astrofísico F. Sánchez 2, 38206 La Laguna, Tenerife, Spain

² Departamento de Química Orgánica, Universidad de La Laguna (ULL), Avda. Astrofísico F. Sánchez 2, 38206 La Laguna, Tenerife, Spain

³ Chemical Engineering Department, University of Almería, 04120 Almería, Spain

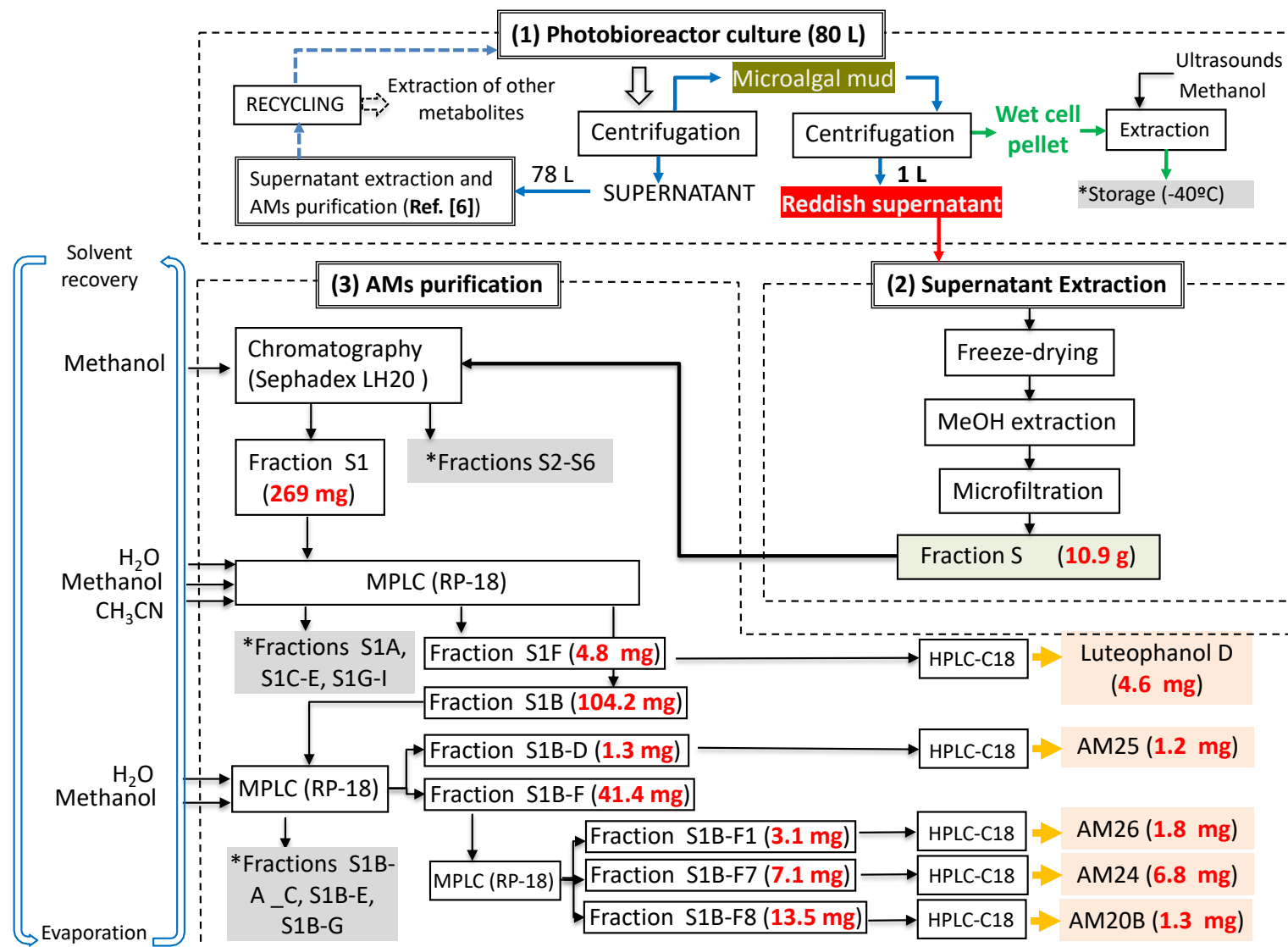
⁴ Research Center CIAIMBITAL, University of Almería, 04120 Almería, Spain

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Scheme S1. Production of new amphidinol analogues by the marine microalga *Amphidinium carterae* grown in a pilot-scale LED-illuminated photobioreactor.



Scheme S2. Isolation procedure for new amphidinol analogues.

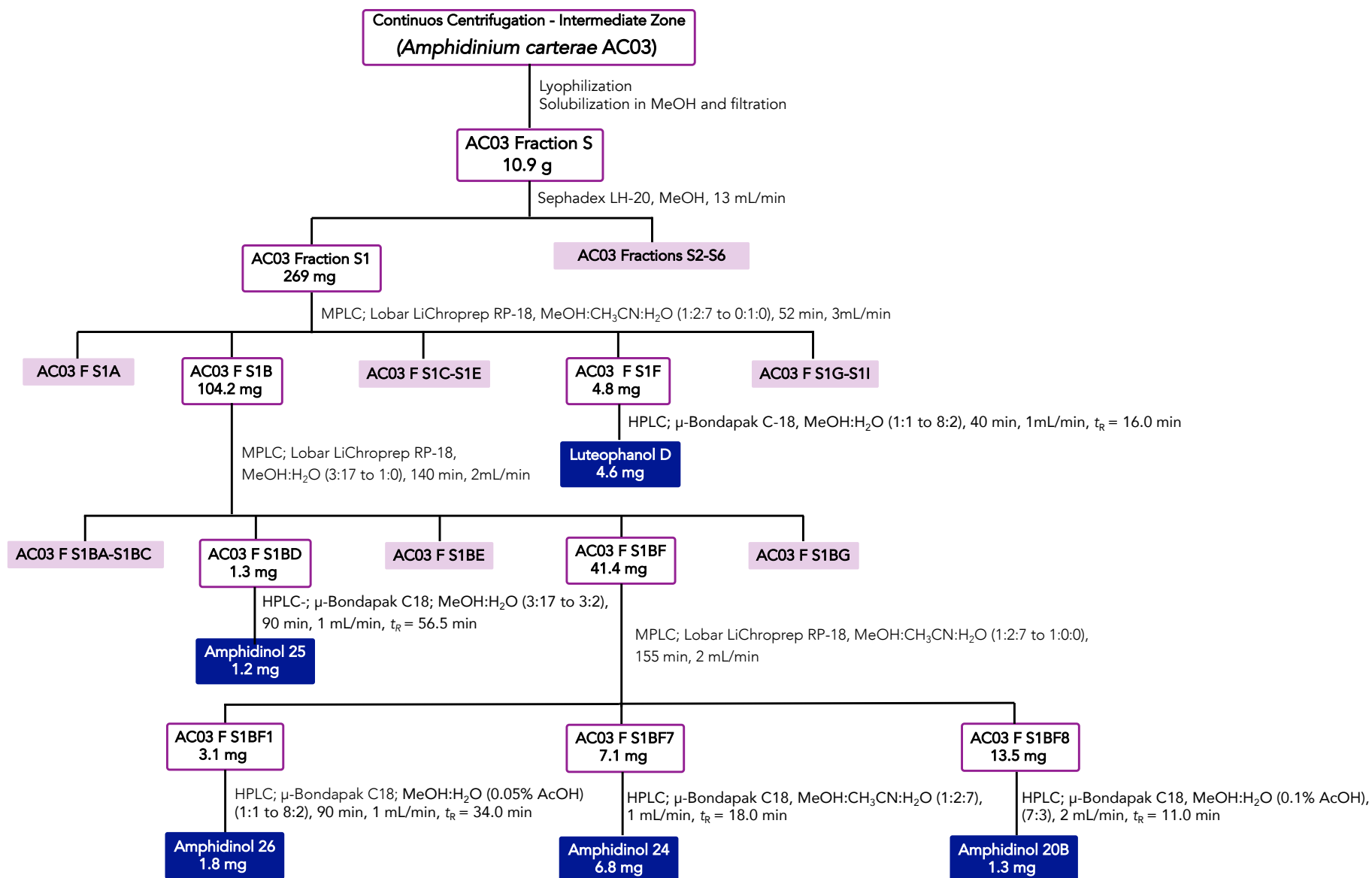


Table S1. ^1H and ^{13}C -NMR data (600 MHz, CD_3OD) for amphidinols 24, 25 and 27.

#		Amphidinol 24		Amphidinol 25		Amphidinol 27		#		Amphidinol 24		Amphidinol 25		Amphidinol 27							
		δC , type	δH	δC , type	δH	δC , type	δH			δC , type	δH	δC , type	δH	δC , type	δH						
1	67.0, CH ₂	3.43; 3.48	67.1, CH ₂	3.43; 3.47	67.1, CH ₂	3.43; 3.48		25	71.1, CH	3.86	71.1, CH	3.86	70.1, CH	3.87	53	37.6, CH ₂	1.60; 1.64	37.6, CH ₂	1.62 (2H)	38.8, CH	2.16 (2H)
2	73.0, CH	3.58	73.1, CH	3.59	73.1, CH	3.59		26	36.2, CH ₂	1.59; 1.68	37.4, CH ₂	1.59; 1.68	36.2, CH ₂	1.59; 1.68	54	72.2, CH	4.12	72.4, CH	4.11	182.8, C	
3	34.2, CH ₂	1.38; 1.54	34.3, CH ₂	1.37; 1.54	34.3, CH ₂	1.38; 1.54		27	36.8, CH ₂	2.12; 2.21	36.5, CH ₂	2.12; 2.21	36.4, CH ₂	1.54; 1.90	55	137.0, CH	5.69	133.8, CH	5.67	6.7, CH ₃	0.97
4	22.6, CH ₂	1.38; 1.62	22.6, CH ₂	1.38; 1.61	22.6, CH ₂	1.38; 1.61		28	139.0, C		139.0, C		139.1, C		56	130.7, CH	6.23	130.7, CH	6.23	17.1, CH ₃	1.75
5	38.2, CH ₂	1.40; 1.50	38.2, CH ₂	1.40; 1.50	38.1, CH ₂	1.40; 1.50		29	125.9, CH	5.48	125.9, CH	5.48	125.8, CH	5.48	57	130.7, CH	6.23	130.7, CH	6.23	112.6, CH ₂	4.99; 5.09
6	72.0, CH	3.54	72.1, CH	3.54	72.0, CH	3.56		30	67.6, CH	4.55	67.6, CH	4.55	67.6, CH	4.56	58	137.0, CH	5.69	133.8, CH	5.67		
7	38.2, CH ₂	1.40; 1.50	38.2, CH ₂	1.40; 1.50	38.1, CH ₂	1.40; 1.50		31	72.0, CH	3.69	72.0, CH	3.69	72.0, CH	3.68	59	72.8, CH	4.10	72.4, CH	4.11		
8	22.6, CH ₂	1.38; 1.62	22.6, CH ₂	1.38; 1.61	22.6, CH ₂	1.38; 1.62		32	78.8, CH	3.96	78.9, CH	3.97	78.8, CH	3.96	60	34.2, CH ₂	1.59; 1.71	33.7, CH ₂	1.71; 1.73		
9	37.6, CH ₂	1.40; 1.52	37.6, CH ₂	1.39; 1.52	37.7, CH ₂	1.40; 1.52		33	67.1, CH	3.97	68.4, CH	4.04	68.4, CH	4.05	61	34.2, CH ₂	1.38; 1.54	34.2, CH ₂	1.71; 1.87		
10	71.9, CH	3.58	72.2, CH	3.58	72.4, CH	3.59		34	68.4, CH	4.04	68.4, CH	3.97	67.1, CH	3.98	62	73.0, CH	3.58	77.3, CH	4.50		
11	41.2, CH ₂	2.20 (2H)	41.4, CH ₂	2.20 (2H)	41.2, CH ₂	2.19 (2H)		35	30.0, CH ₂	1.79 (2H)	30.1, CH ₂	1.79 (2H)	30.1, CH ₂	1.79 (2H)	63	67.8, CH ₂	3.43; 3.48	69.1, CH ₂	4.10; 4.26		
12	128.6, CH	5.69	128.6, CH	5.68	128.5, CH	5.70		36	75.3, CH	3.49	75.3, CH	3.49	75.3, CH	3.49	64	6.6, CH ₃	0.98	6.7, CH ₃	0.98		
13	136.0, CH	5.53	135.9, CH	5.53	135.9, CH	5.55		37	74.2, CH	3.60	74.1, CH	3.60	74.1, CH	3.61	65	17.1, CH ₃	1.75	17.1, CH ₃	1.75		
14	73.2, CH	4.05	73.3, CH	4.05	73.2, CH	4.05		38	32.1, CH ₂	1.57; 1.97	32.3, CH ₂	1.57; 1.97	32.2, CH ₂	1.56; 1.97	66	112.8, CH ₂	4.99; 5.08	112.7, CH ₂	4.99; 5.08		
15	41.7, CH ₂	2.25 (2H)	41.8, CH ₂	2.24 (2H)	41.7, CH ₂	2.24 (2H)		39	27.8, CH ₂	2.10; 2.42	27.9, CH ₂	2.10; 2.42	28.0, CH ₂	2.10; 2.41							
16	129.7, CH	5.54	129.6, CH	5.53	129.6, CH	5.55		40	151.4, C		151.1, C		151.2, C								
17	137.3, CH	5.60	130.1, CH	5.60	130.1, CH	5.60		41	76.3, CH	4.18	76.2, CH	4.18	76.1, CH	4.19							
18	37.7, CH ₂	2.08; 2.48	37.7, CH ₂	2.08; 2.48	37.7, CH ₂	2.08; 2.48		42	74.1, CH	3.35	75.0, CH	3.34	75.0, CH	3.35							
19	72.2, CH	3.52	72.2, CH	3.52	72.1, CH	3.52		43	70.0, CH	4.05	70.1, CH	4.04	70.2, CH	4.04							
20	78.9, CH	3.52	78.7, CH	3.52	78.7, CH	3.52		44	31.1, CH ₂	1.56; 2.09	31.3, CH ₂	1.56; 2.09	31.2, CH ₂	1.56; 2.09							
21	35.0, CH	2.30	35.0, CH	2.30	34.9, CH	2.30		45	66.8, CH	4.05	67.1, CH	4.05	67.2, CH	4.05							
22	79.9, CH	3.53	79.6, CH	3.53	79.7, CH	3.53		46	68.4, CH	4.05	68.4, CH	4.04	68.4, CH	4.05							
23	71.7, CH	3.71	71.2, CH	3.71	71.7, CH	3.72		47	80.2, CH	3.74	80.3, CH	3.75	80.1, CH	3.75							
24	40.7, CH ₂	1.54; 1.91	40.9, CH ₂	1.53; 1.91	40.8, CH ₂	1.54; 1.90		48	71.6, CH	3.97	71.7, CH	3.96	71.6, CH	3.97							
								49	73.8, CH	4.37	73.9, CH	4.36	73.7, CH	4.37							
								50	128.6, CH	5.64	128.6, CH	5.63	128.5, CH	5.66							
								51	134.9, CH	5.80	135.0, CH	5.80	134.7, CH	5.83							
								52	29.3, CH ₂	2.16 (2H)	29.4, CH ₂	2.15 (2H)	29.4, CH ₂	2.18 (2H)							

[illegible]

Figure S2. COSY spectrum (600 MHz, CD₃OD) for amphidinol 24.

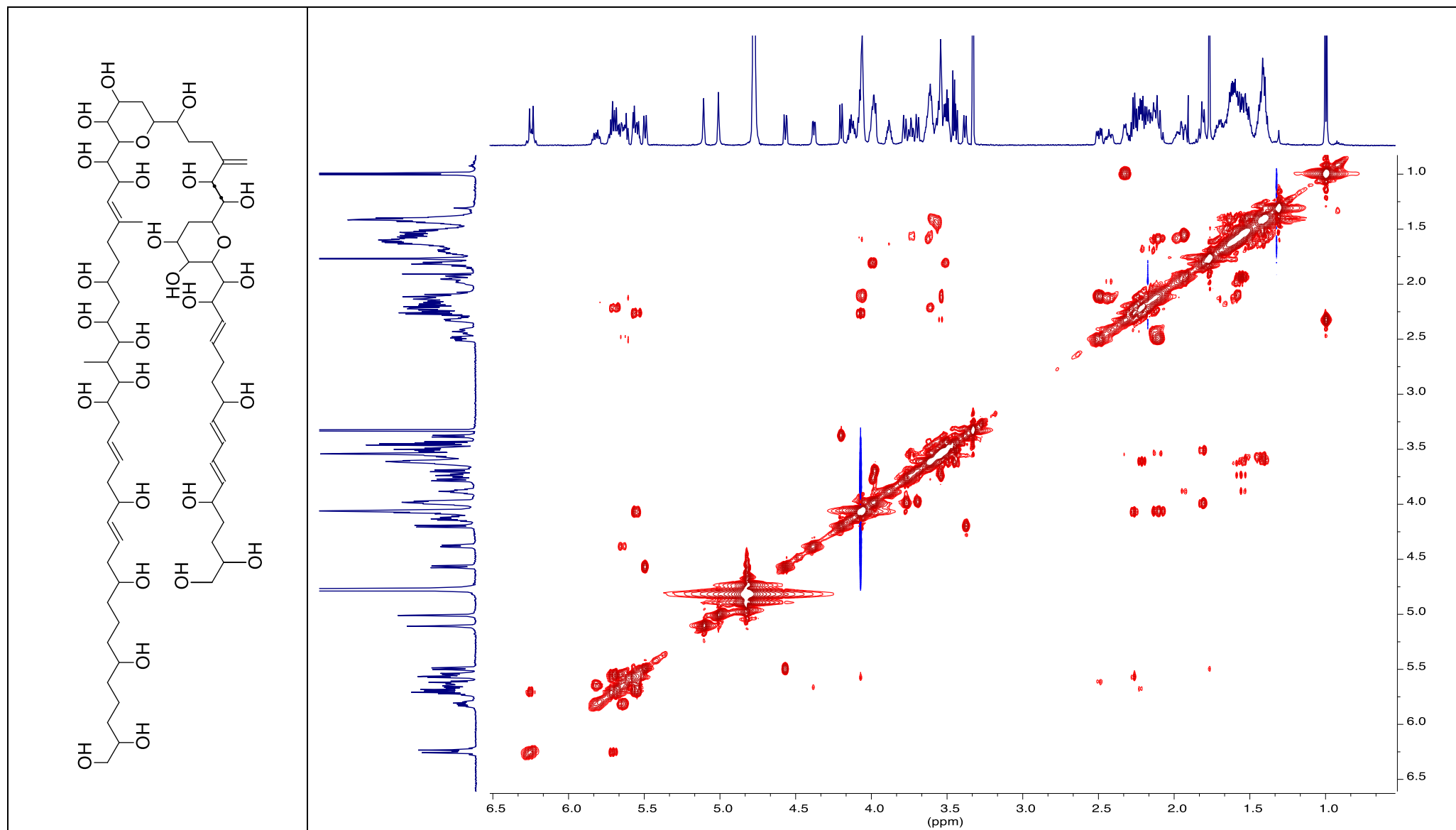


Figure S3. HSQC_{ed} spectrum (600 MHz, CD₃OD) for amphidinol 24.

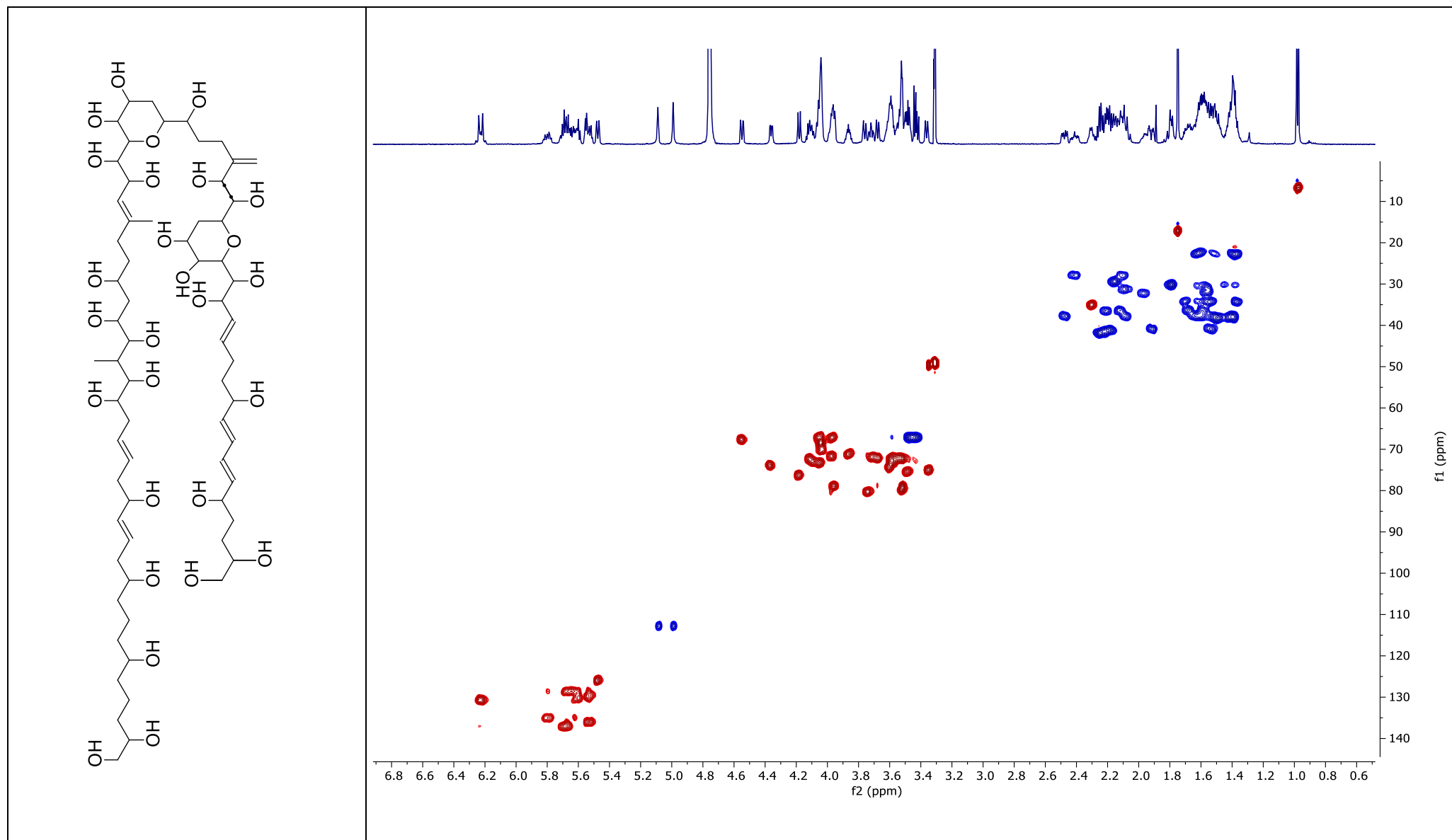
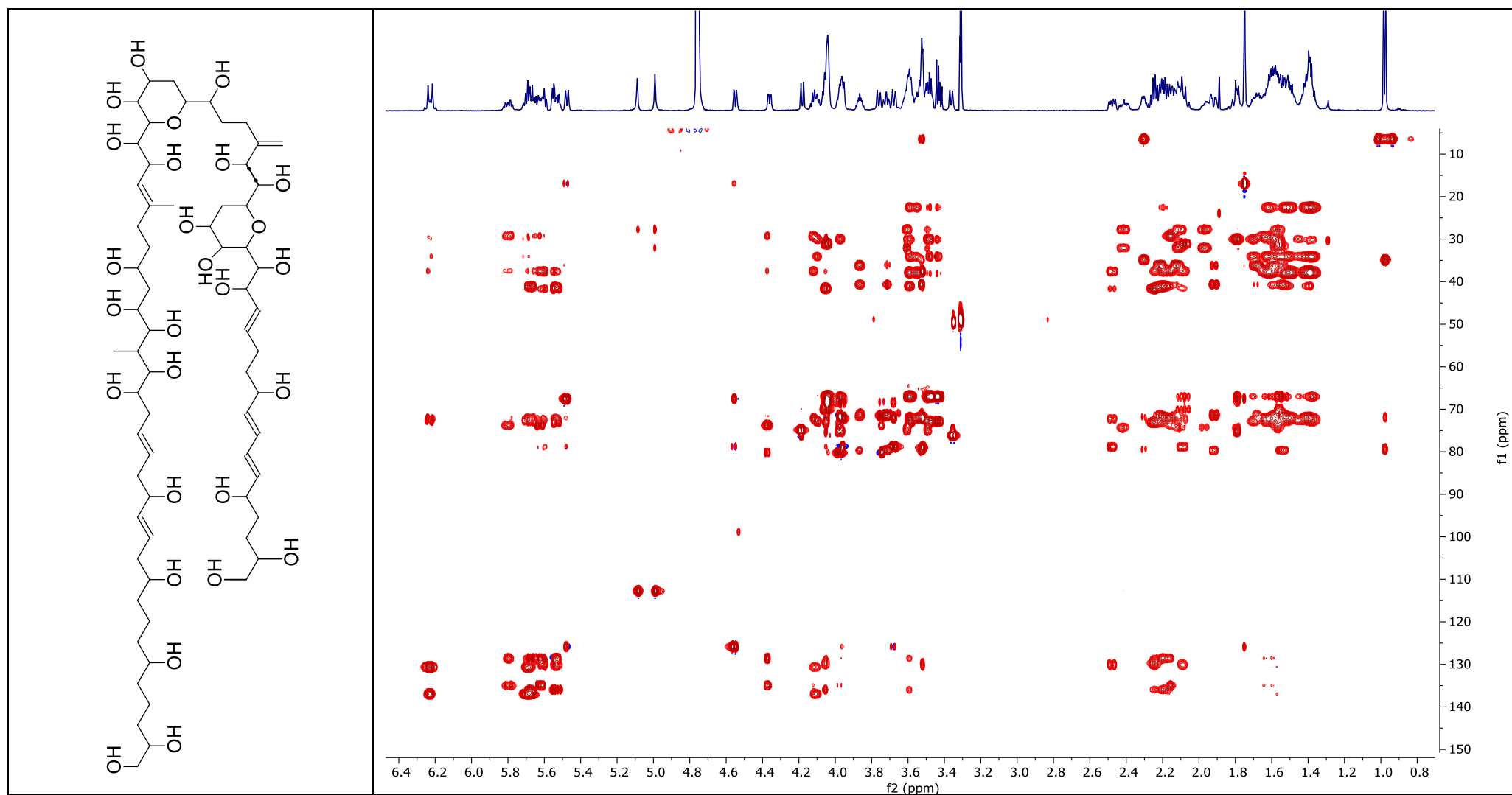


Figure S4. HSQC-TOCSY spectrum (600 MHz, CD₃OD) for amphidinol 24.



The figure displays the chemical structure of a complex polyhydroxy compound and its corresponding NMR spectra. The chemical structure is a long, branched molecule with multiple hydroxyl groups and a central ether linkage. The NMR spectra include a 1D ¹H NMR spectrum at the top and a 2D COSY spectrum below it. The 1D spectrum shows peaks in the aromatic region (6.5-7.5 ppm), a sugar region (4.5-6.0 ppm), and a long aliphatic chain region (1.0-2.5 ppm). The COSY spectrum shows correlations between protons in the aliphatic chain and the sugar moiety.

Figure S6. H2BC spectrum(600 MHz, CD₃OD) for amphidinol 24.

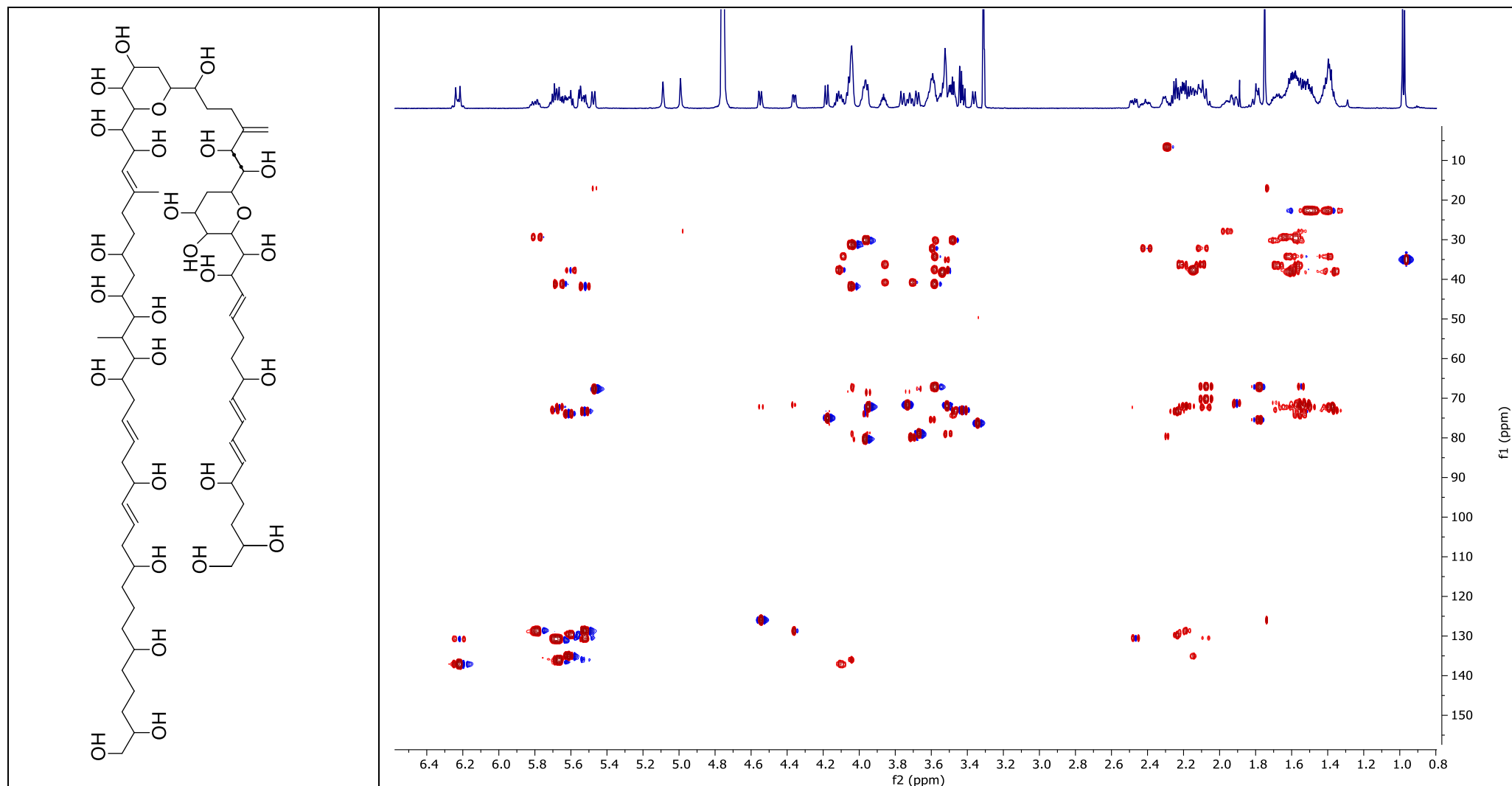


Figure S7. T-ROESY spectrum (600 MHz, CD₃OD) for amphidinol 24.

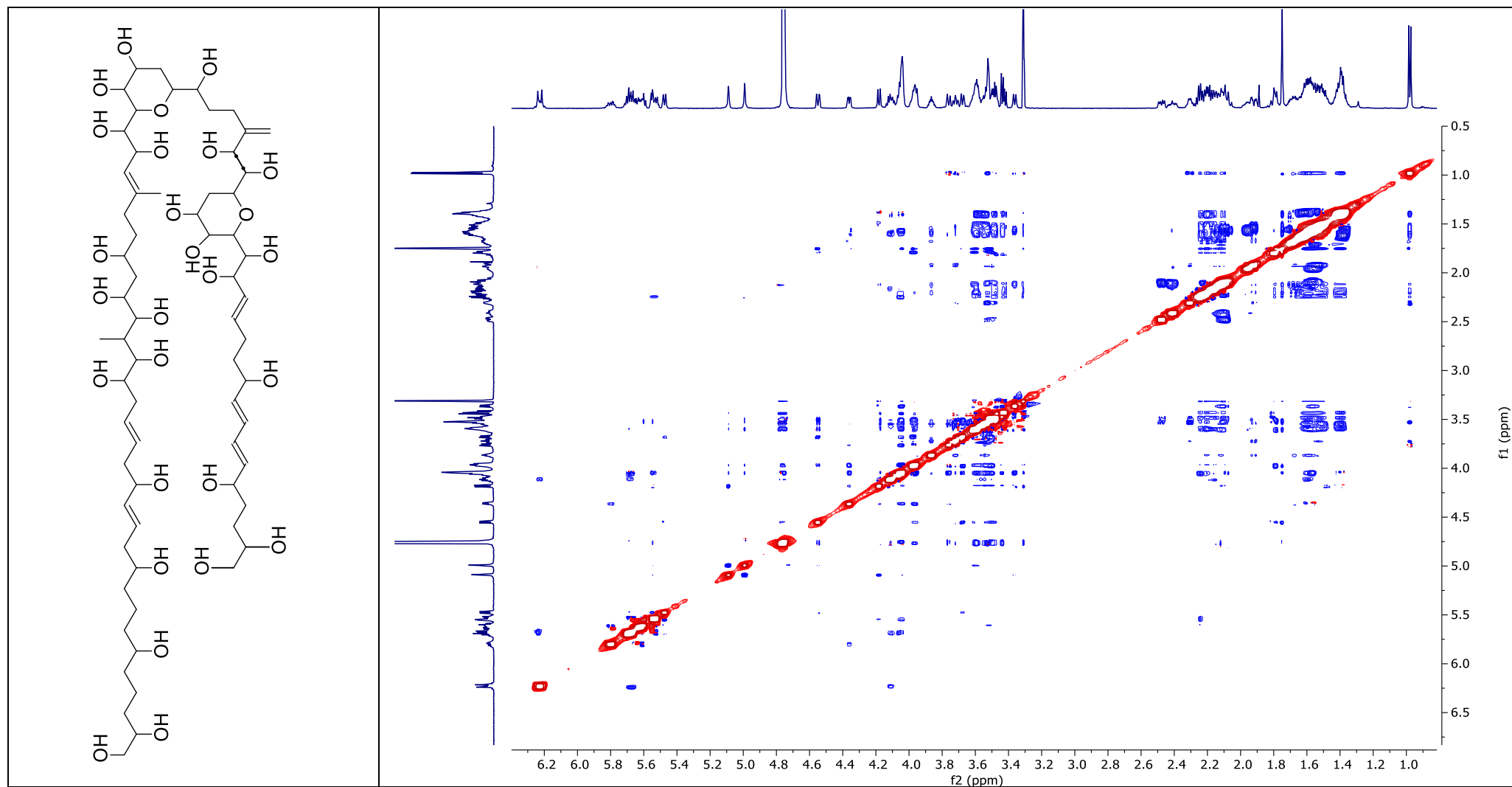


Figure S8. 1D-NOESY spectra (600 MHz, CD₃OD) for amphidinol 24.

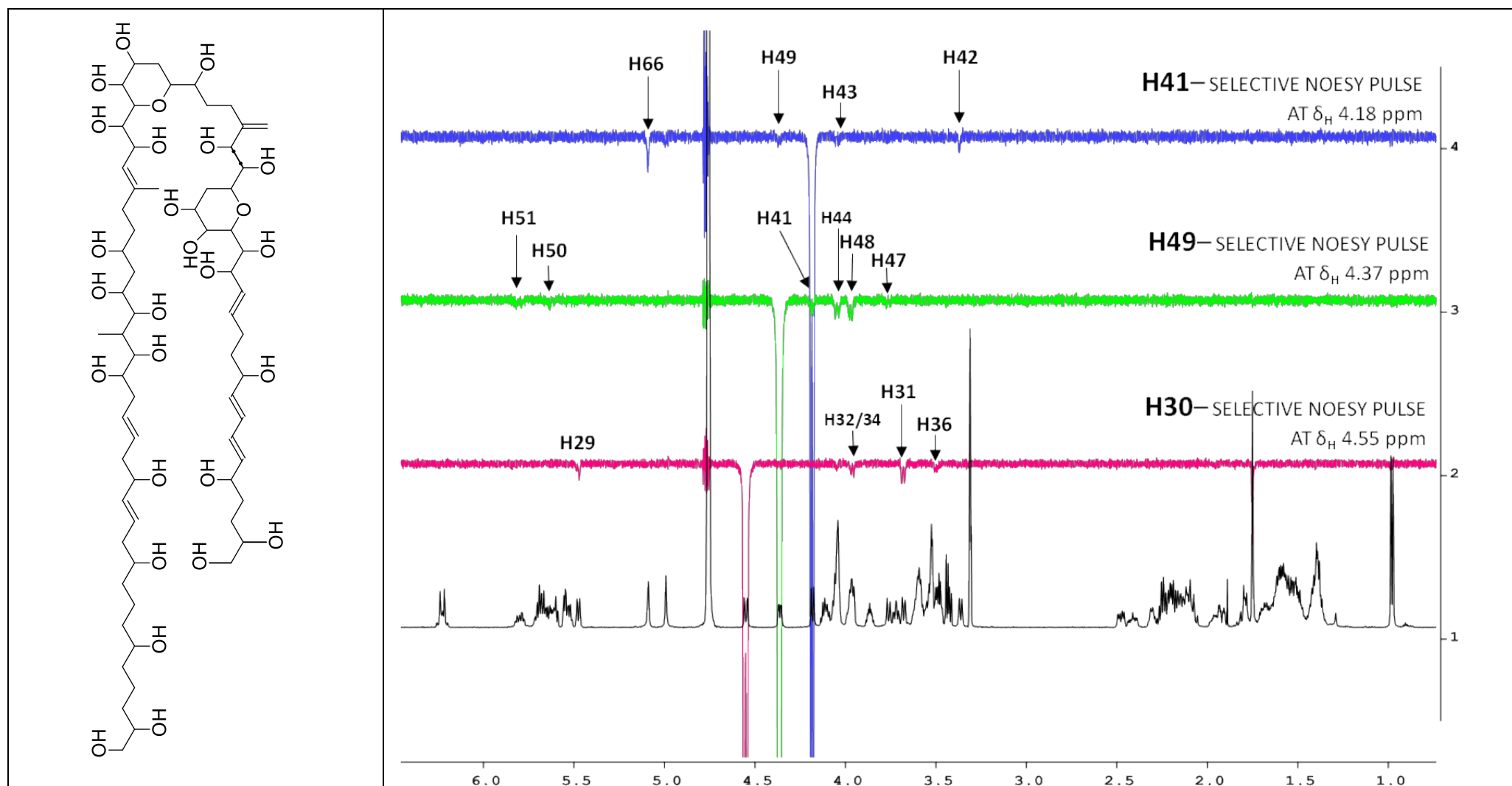


Figure S9. ^1H NMR spectrum (600 MHz, $\text{CD}_3\text{OD}-\text{C}_5\text{D}_5\text{N}$ 2:1) for amphidinol 24.

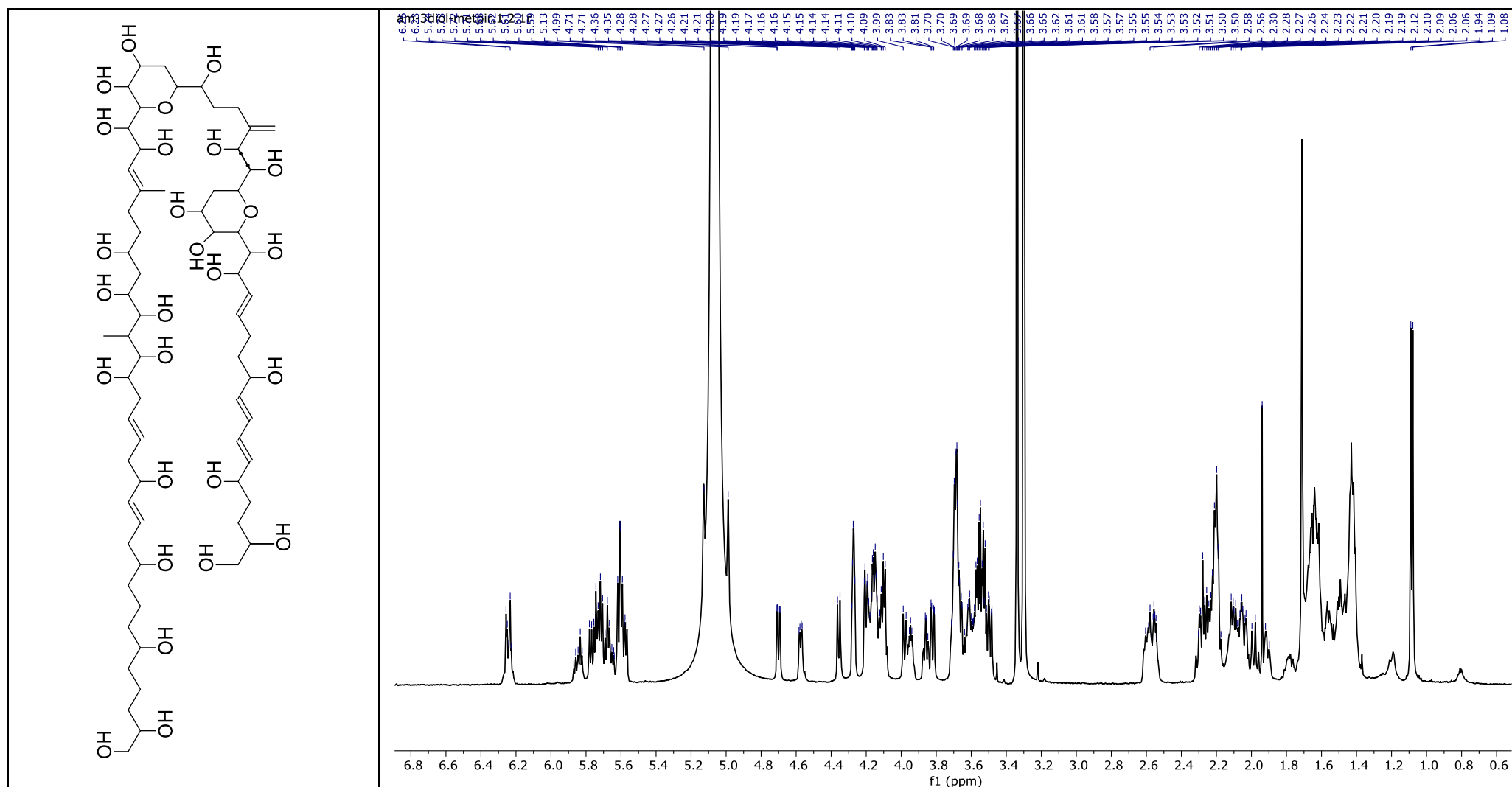


Figure S10. HSQC_{ed} spectrum (600 MHz, CD₃OD-C₅D₅N 2:1) for amphidinol 24.

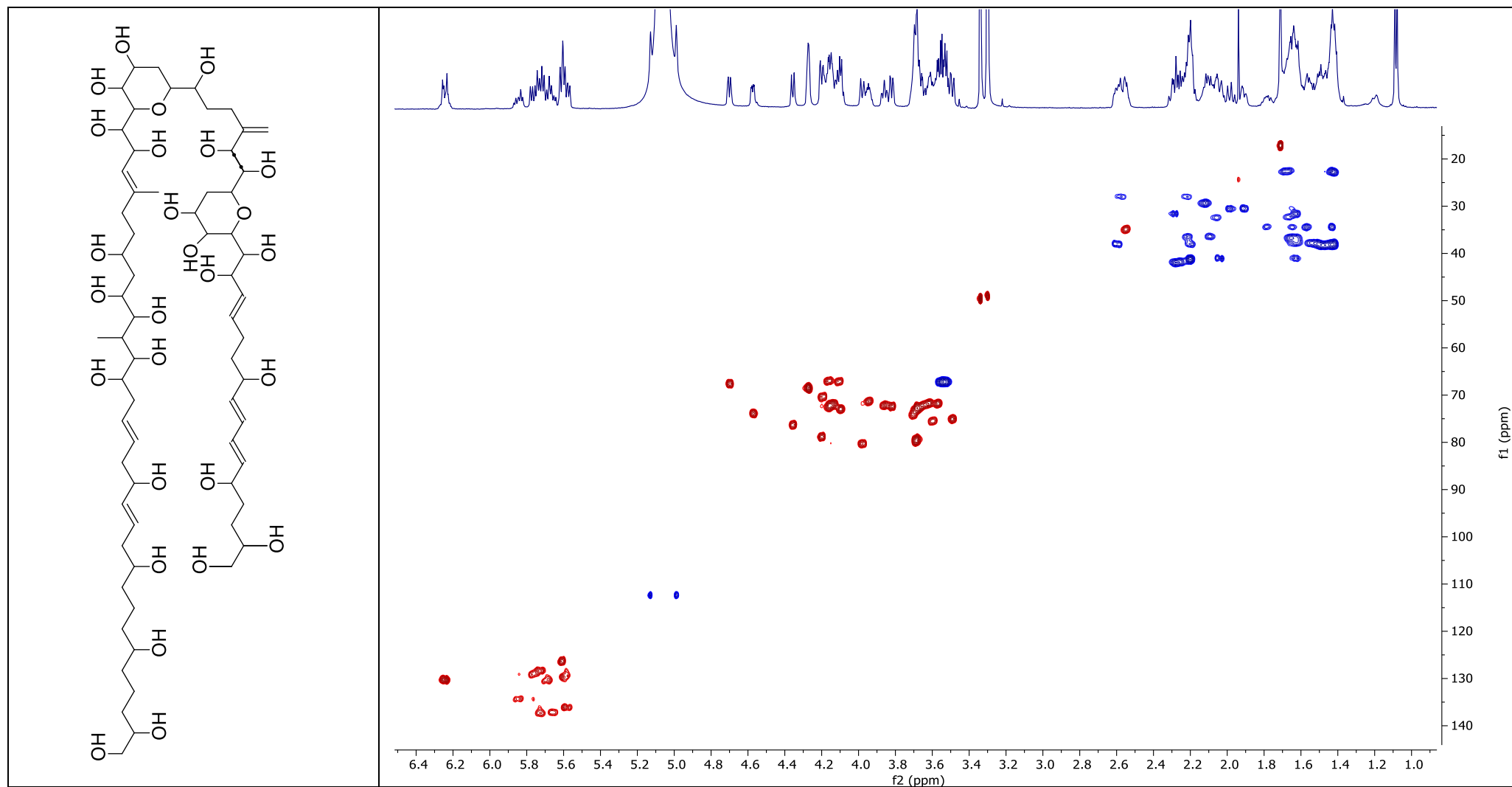


Figure S11. HSQC-TOCSY spectrum (600 MHz, CD₃OD-C₅D₅N 2:1) for amphidinol 24.

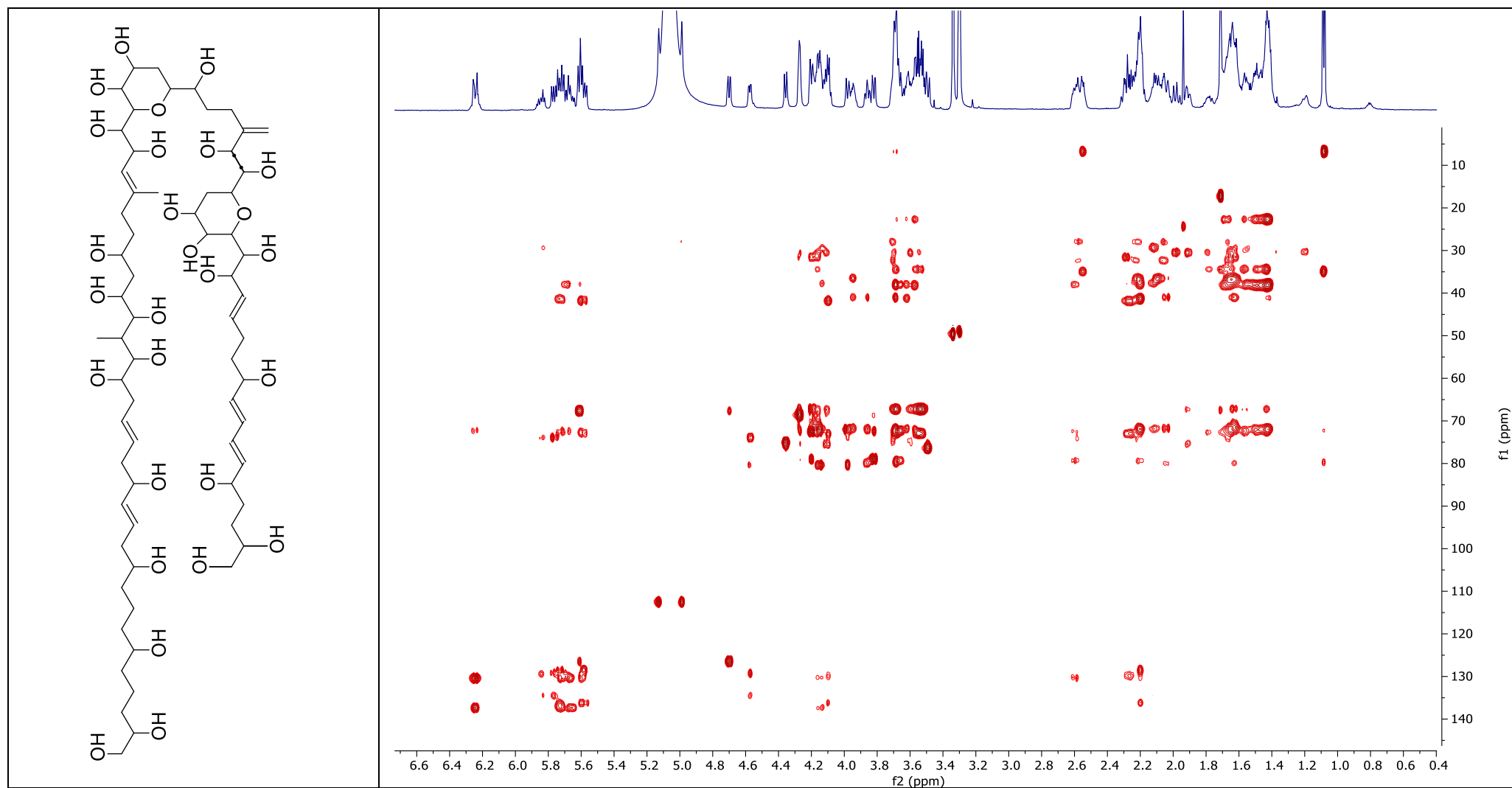


Figure S12. HMBC spectrum (600 MHz, CD₃OD-C₅D₅N 2:1) for amphidinol 24.

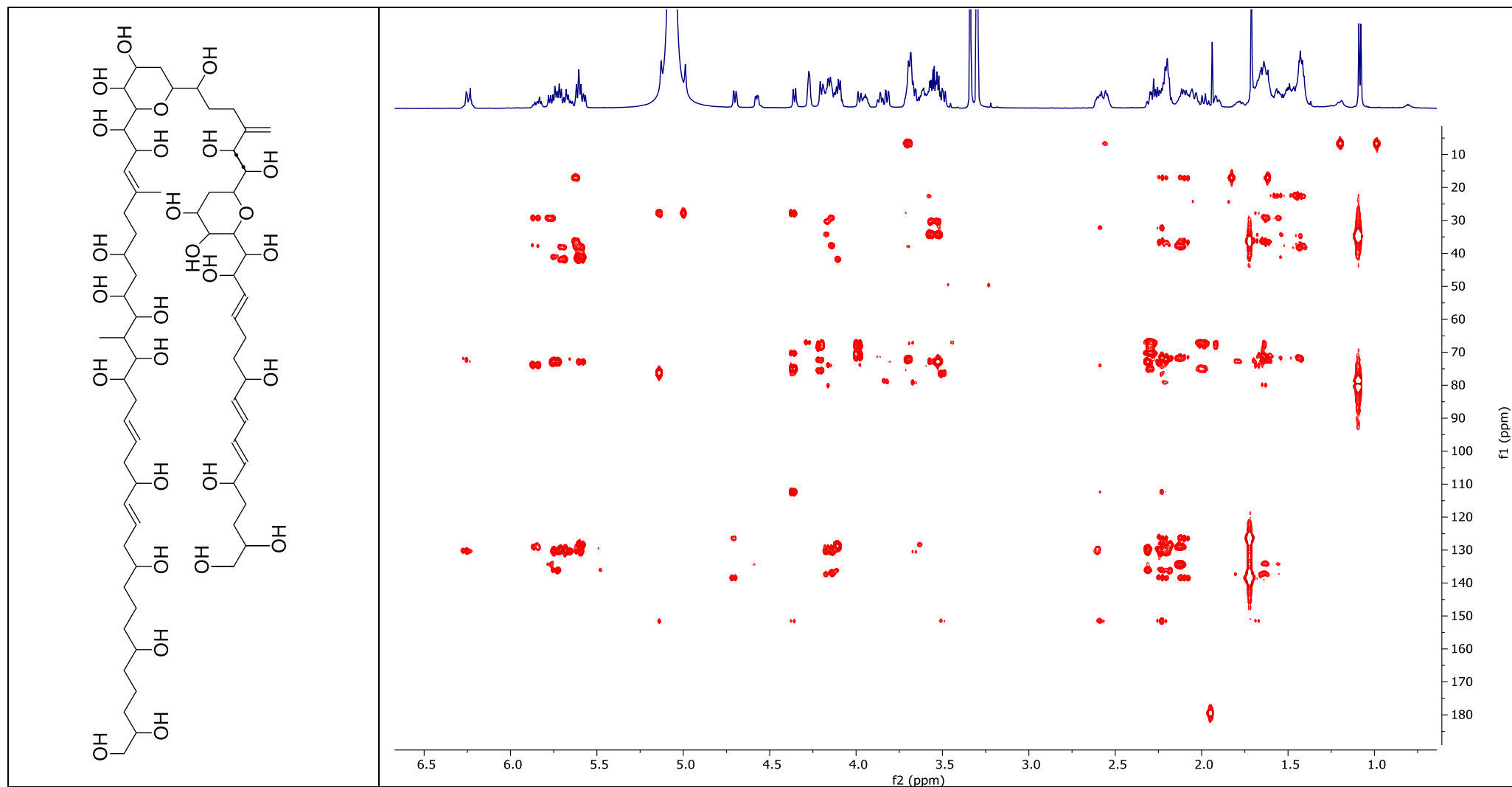


Figure S13. H2BC spectrum (600 MHz, CD₃OD-C₅D₅N 2:1) for amphidinol 24.

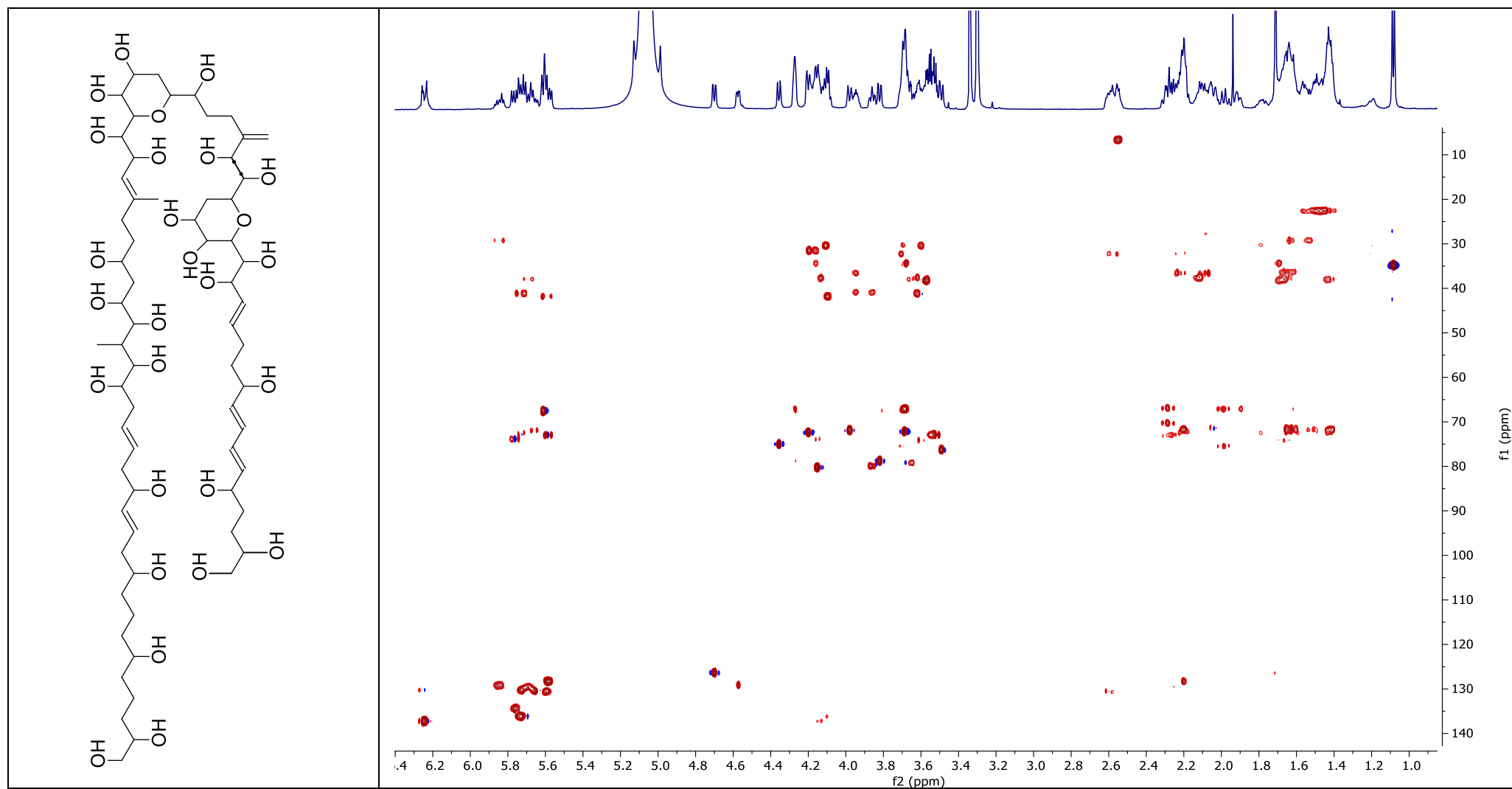


Figure S14. T-ROESY spectrum (600 MHz, CD₃OD-C₅D₅N 2:1) for amphidinol 24.

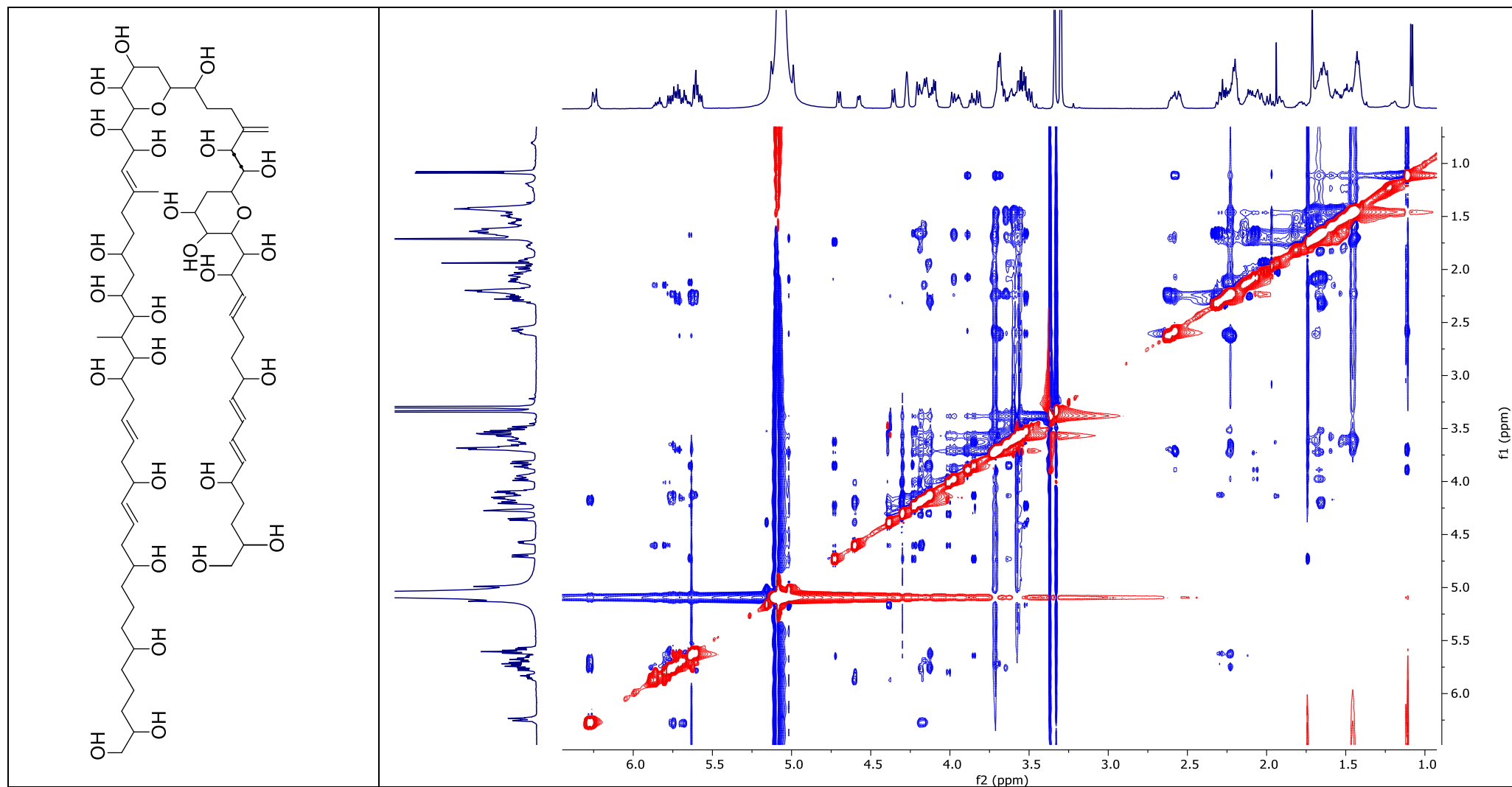


Figure S15. 1D-NOESY spectrum (600 MHz, CD₃OD-C₅D₅N 2:1) for amphidinol 24.

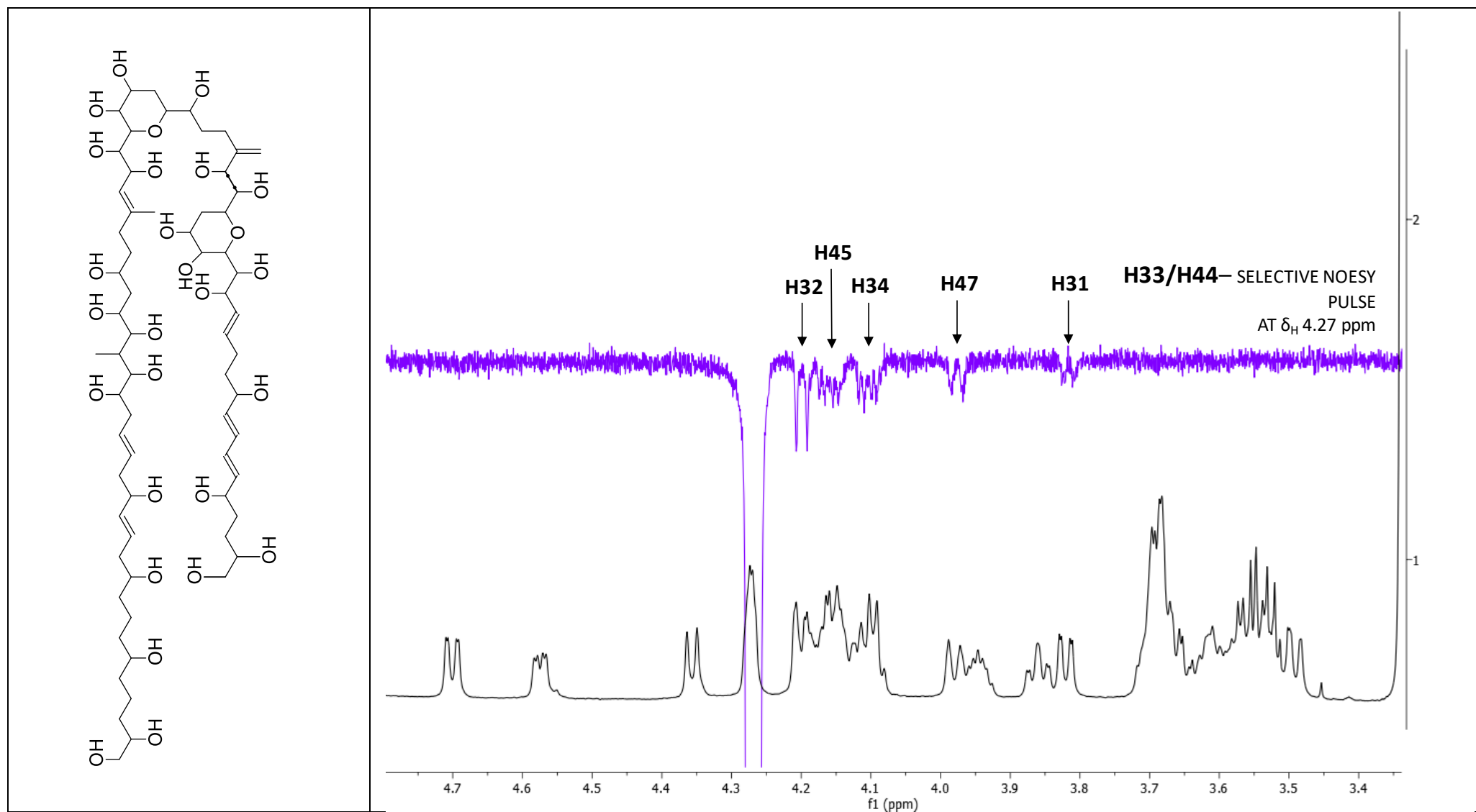


Table S2. ^1H and ^{13}C NMR data comparison for carbons C-30 \rightarrow C-51 in $\text{CD}_3\text{OD}-\text{C}_5\text{D}_5\text{N}$ 2:1 for amphidinol 24 *versus* related synthetic fragments 4a and 4b reported by Wakamiya et al [22].

#	δC , type			#	δH		
	$\text{CD}_3\text{OD}-\text{C}_5\text{D}_5\text{N}$ 2:1	4a	am 24 / 4a		$\text{CD}_3\text{OD}-\text{C}_5\text{D}_5\text{N}$ 2:1	4a	am 24 / 4b
30	67.6, CH	71.1	-3,5	30	4.7	4.08	0,62
31	72.4, CH	68.3	4,1	31	3.82	4.14	-0,32
32	78.9, CH	78.3	0,6	32	4.2	4.26	-0,06
33	68.6, CH	69.1	-0,5	33	4.27	4.33	-0,06
34	67.2, CH	67.3	-0,1	34	4.11	4.11	0
35	30.5, CH_2	31.7	-1,2	35	1.91	1.69	0,22
36	75.5, CH	74.8	0,7	36	1.99	2.08	-0,09
37	74.1, CH	74.2	-0,1	37	3.6	3.62	-0,02
38	32.4, CH_2	32.2	0,2	38	3.7	3.65	0,05
					1.67	1.73	-0,06
39	28.0, CH_2	27.8	0,2	39	2.06	1.91	0,15
					2.22	2.24	-0,02
40	151.5, C	152.0	-0,5	40	2.58	2.54	0,04
41	76.3, CH	76.5	-0,2	41	4.36	4.39	-0,03
42	75.1, CH	75.4	-0,3	42	3.5	3.51	-0,01
43	70.4, CH	70.6	-0,2	43	4.19	4.19	0
					1.63	1.64	-0,01
44	31.6, CH_2	31.9	-0,3	44	2.29	2.34	-0,05
					4.16	4.18	-0,02
45	67.0, CH	67.3	-0,3	45	4.28	4.34	-0,06
46	68.5, CH	68.7	-0,2	46	3.97	4.02	-0,05
47	80.3, CH	80.6	-0,3	47	4.15	4.20	-0,05
48	72.0, CH	72.2	-0,2	48	4.57	4.63	-0,06
49	73.9, CH	74.1	-0,2	49	5.76	5.78	-0,02
50	129.2, CH	129.8	-0,6	50	5.81	5.81	0,04
51	134.4, CH	134.1	0,3	51			

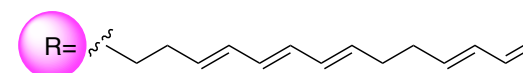
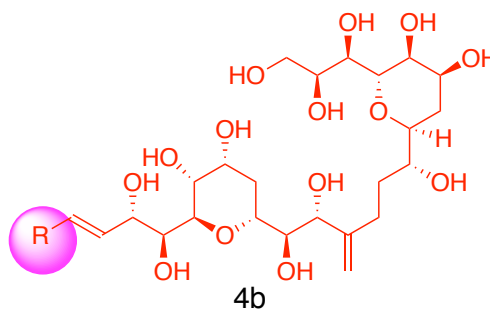
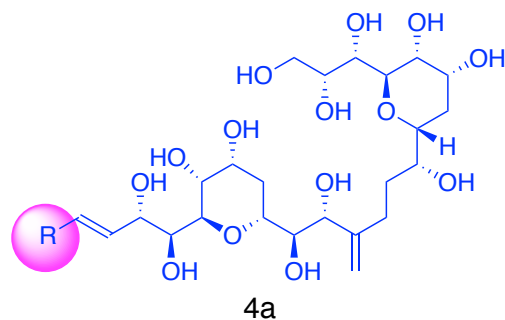
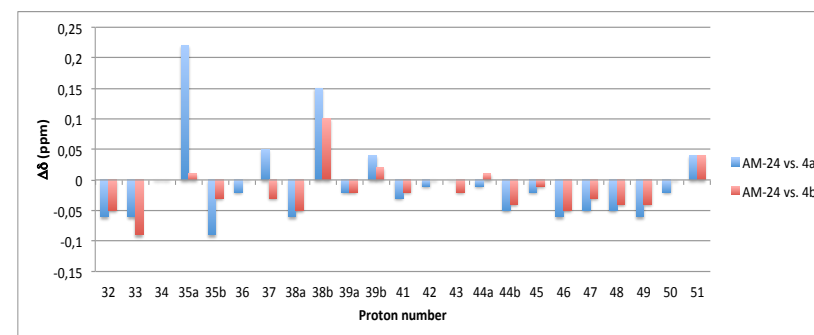
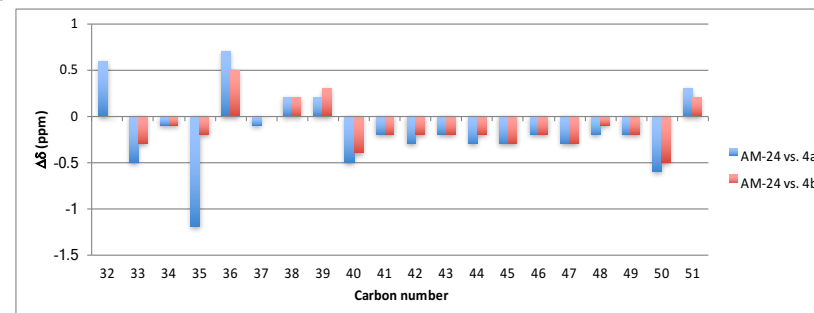


Figure S16. HRESIMS spectrum for amphidinol 24.

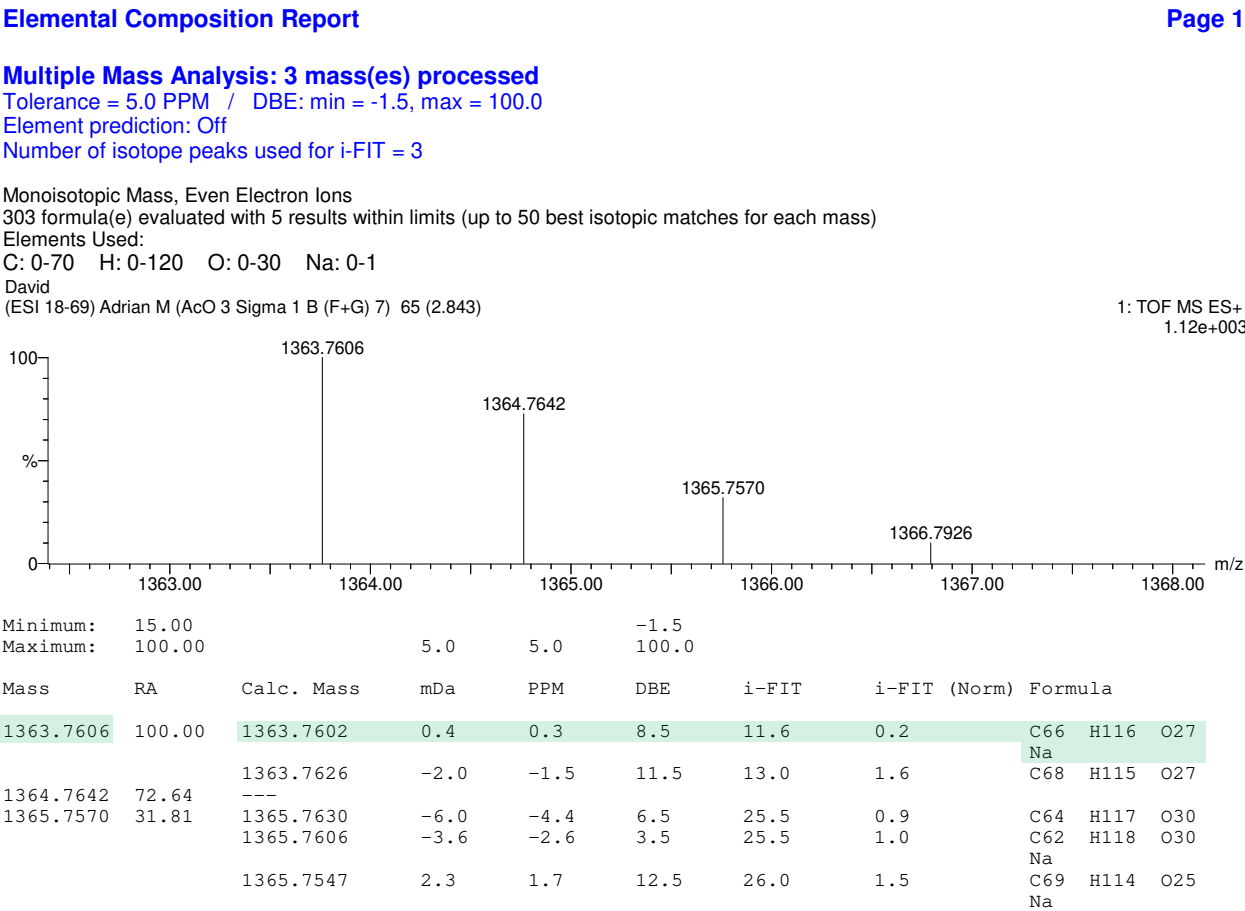


Figure S17. Main MS/MS fragments observed for amphidinol 24.

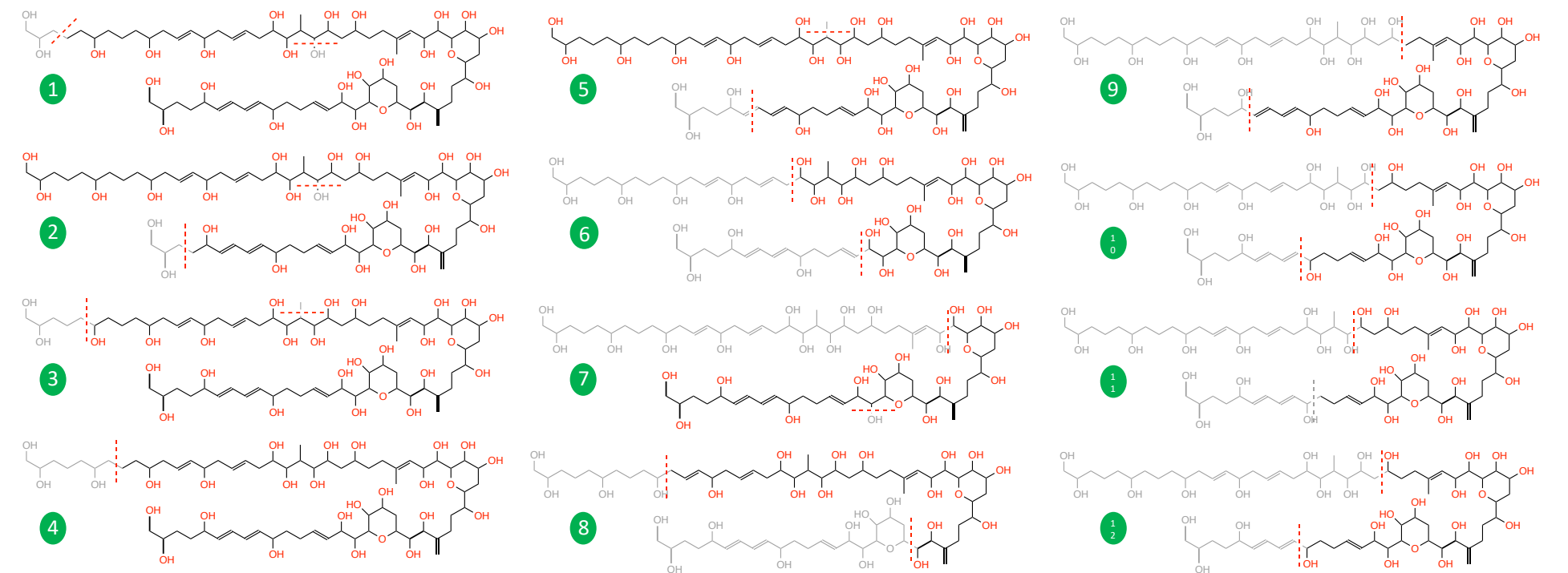
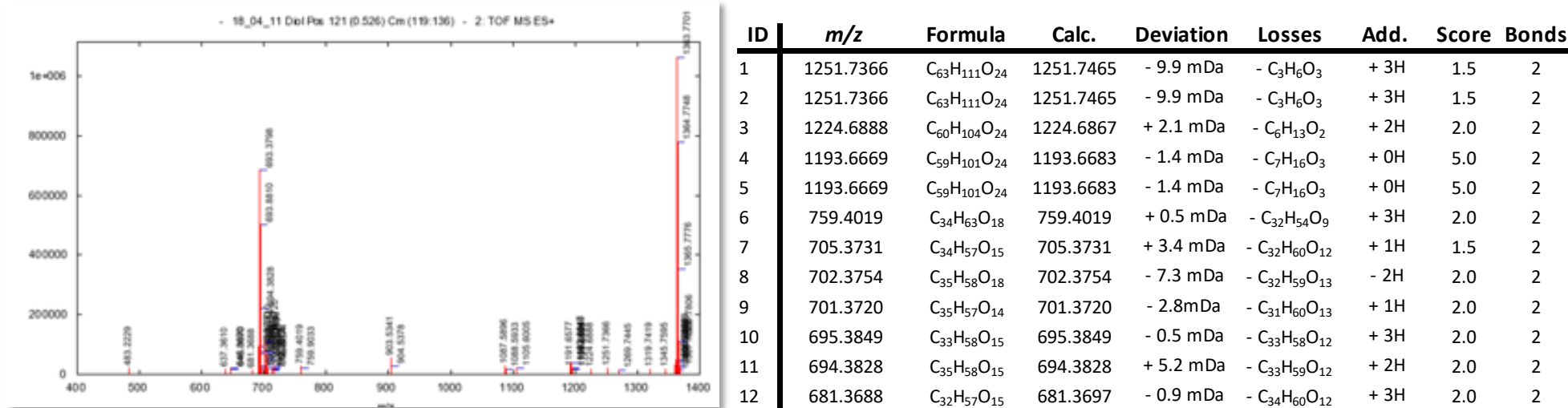


Figure S18. ^1H NMR spectrum (600 MHz, CD_3OD) for amphidinol 25.

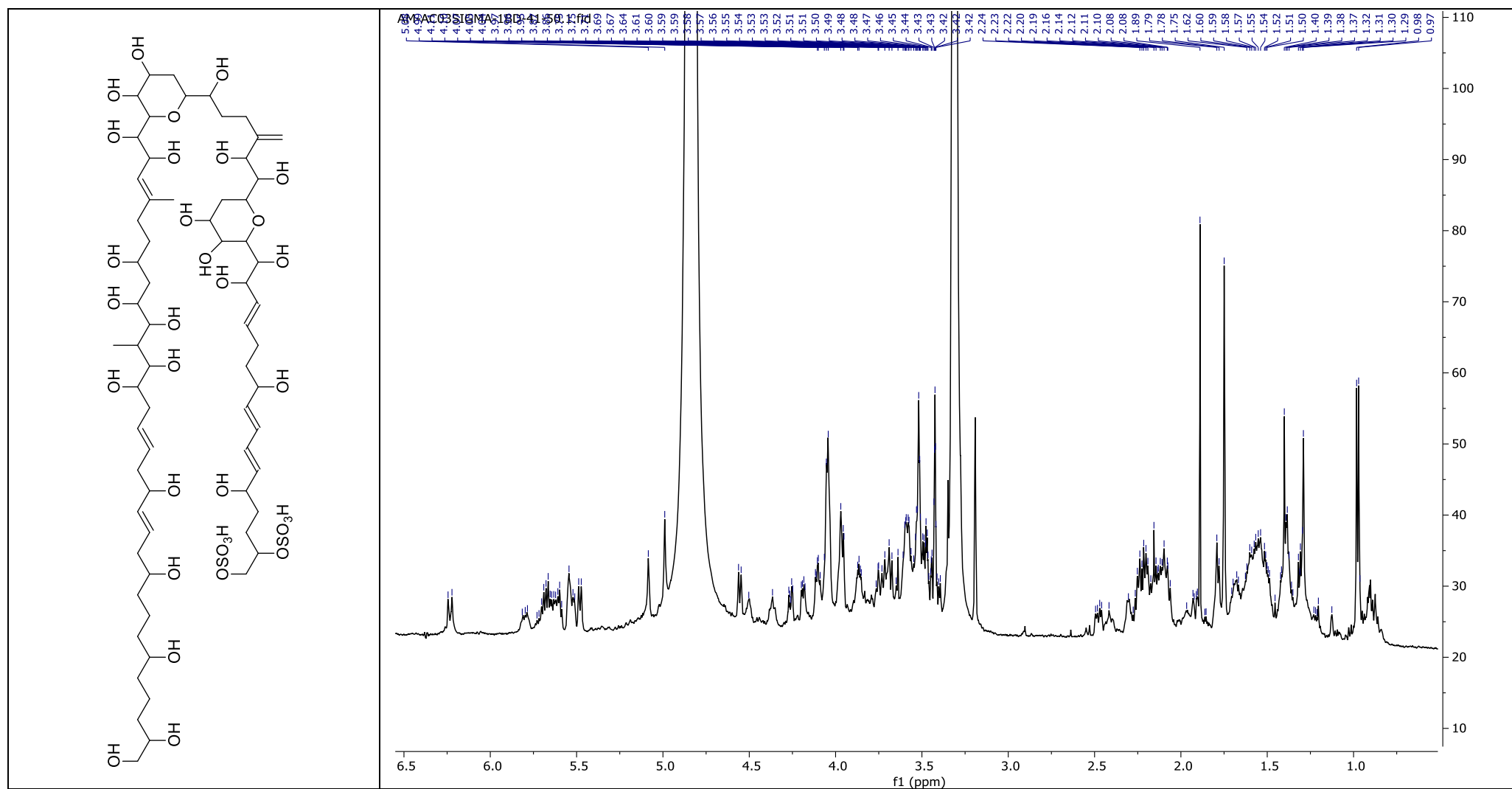


Figure S19. COSY spectrum (600 MHz, CD₃OD) for amphidinol 25.

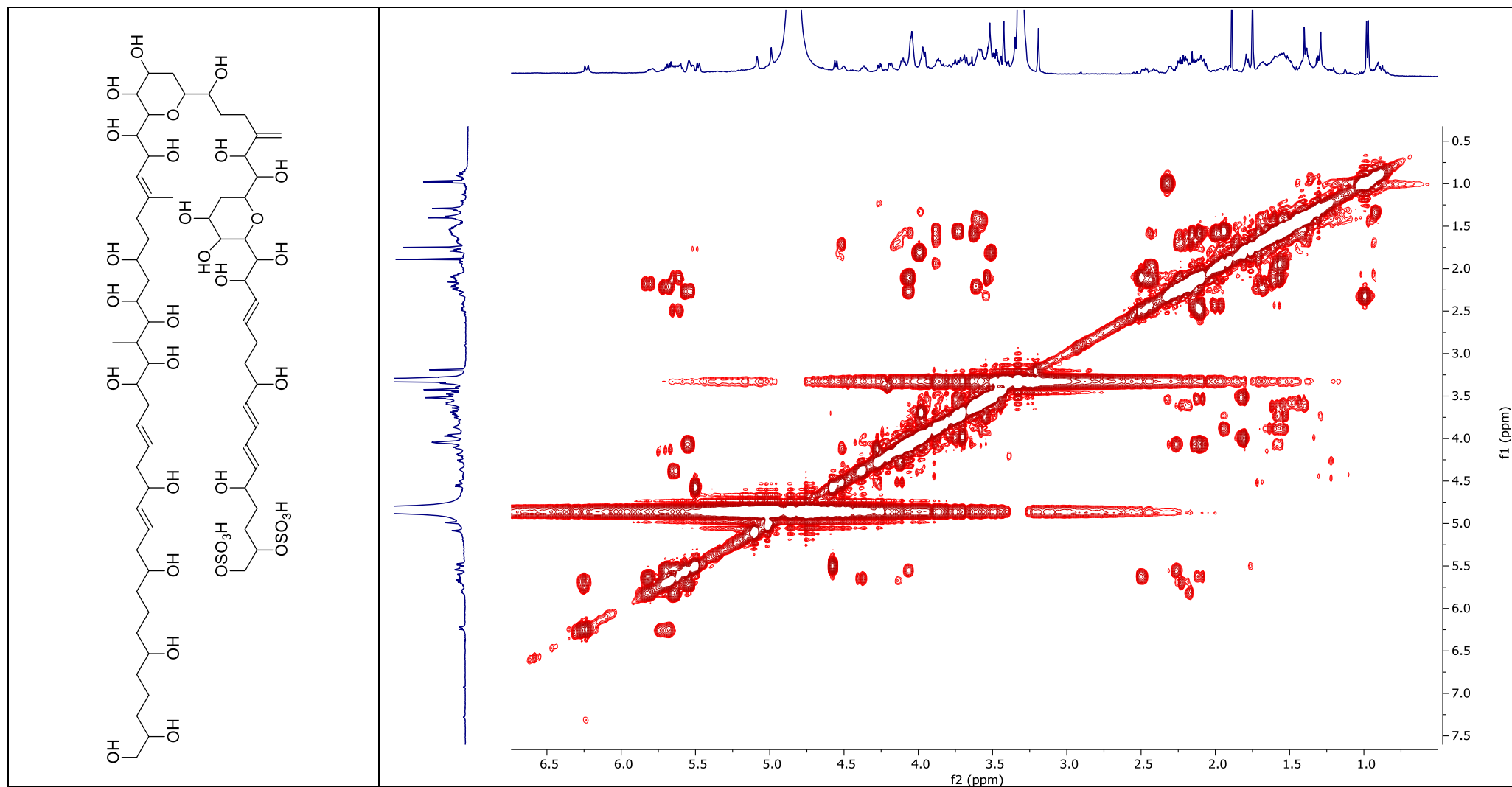


Figure S20. HSQC_{ed} spectrum (600 MHz, CD₃OD) for amphidinol 25.

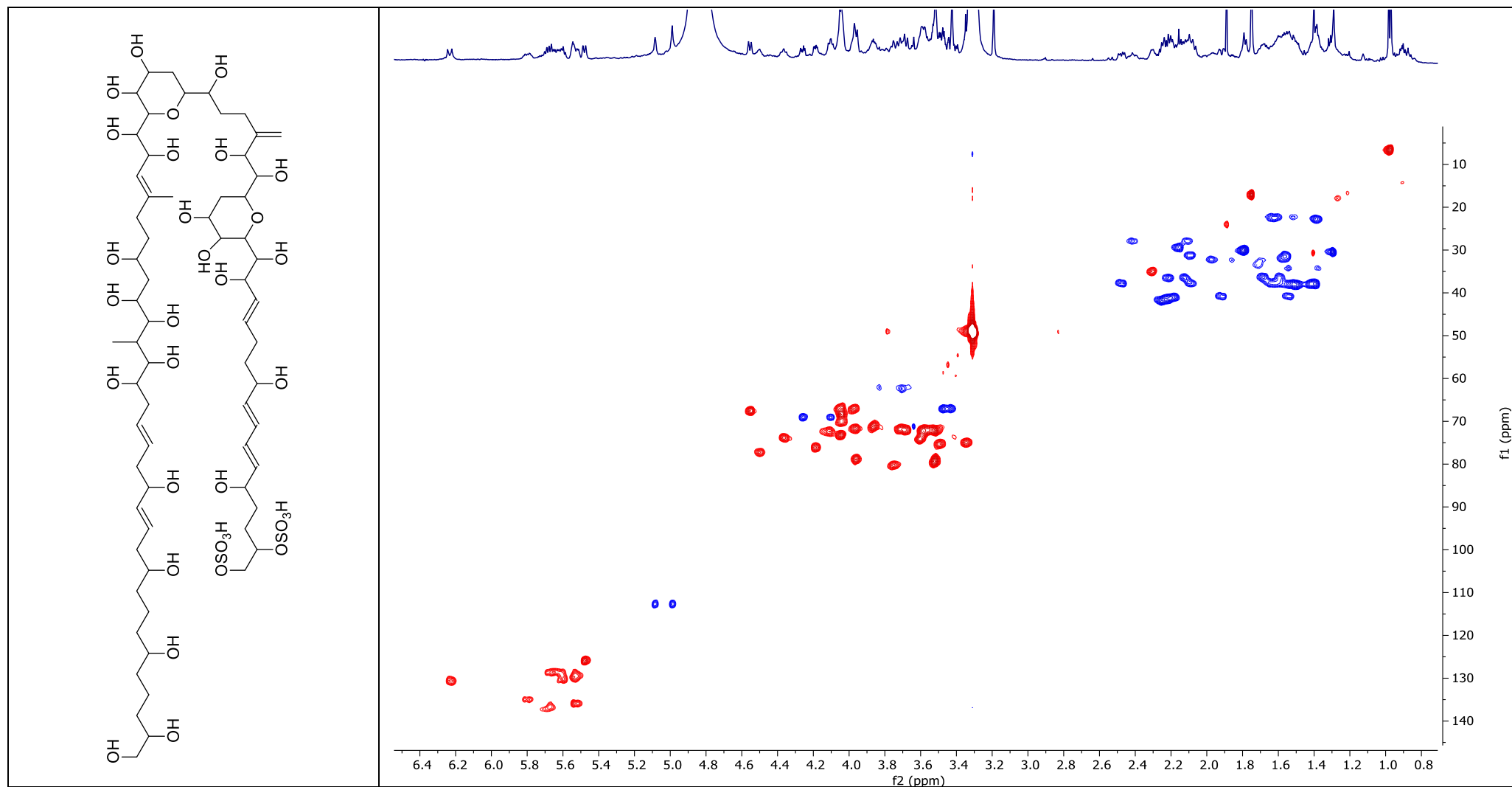


Figure S21. HSQC-TOCSY spectrum (600 MHz, CD₃OD) for amphidinol 25.

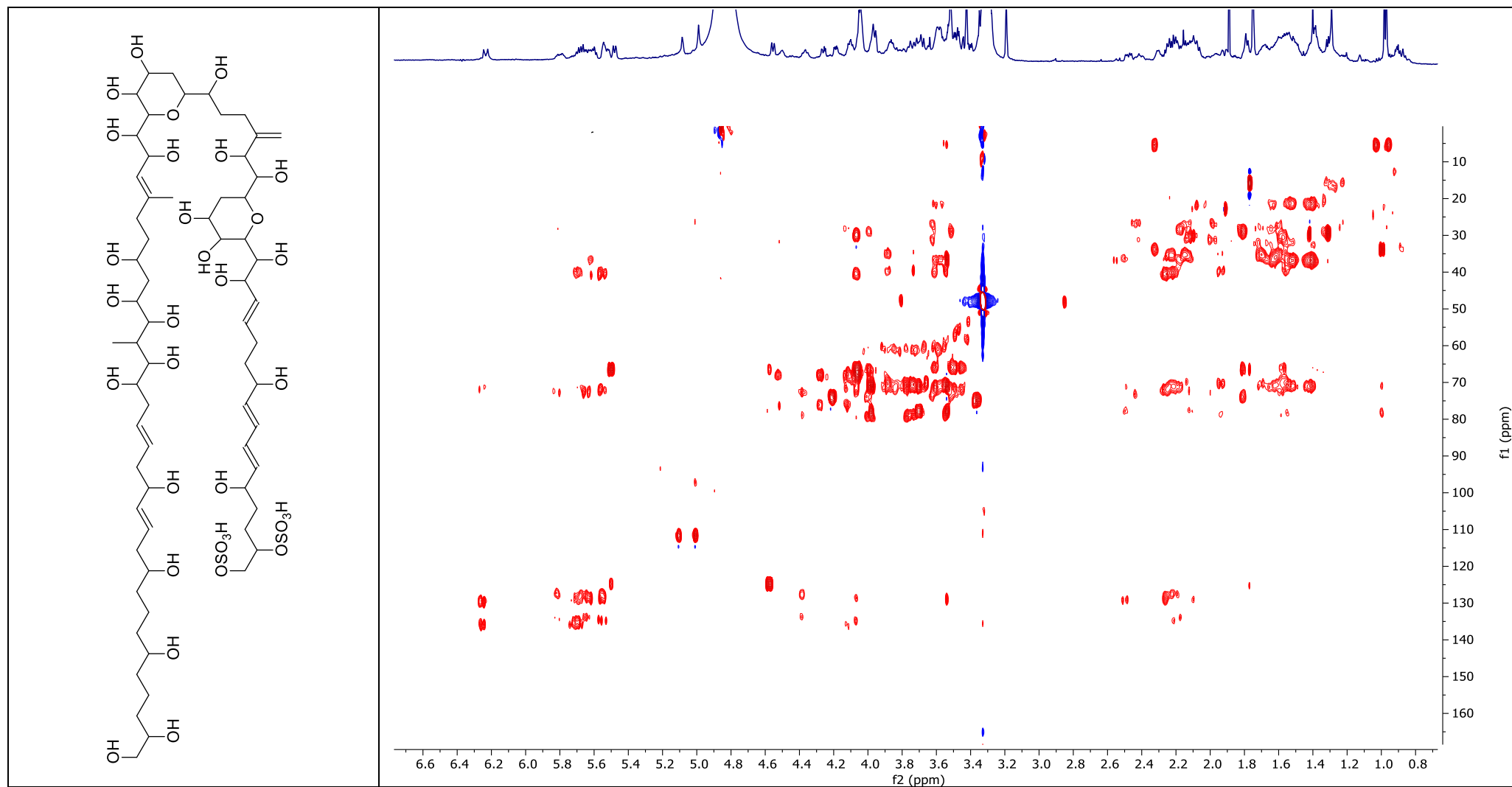


Figure S22. HMBC spectrum (600 MHz, CD₃OD) for amphidinol 25.

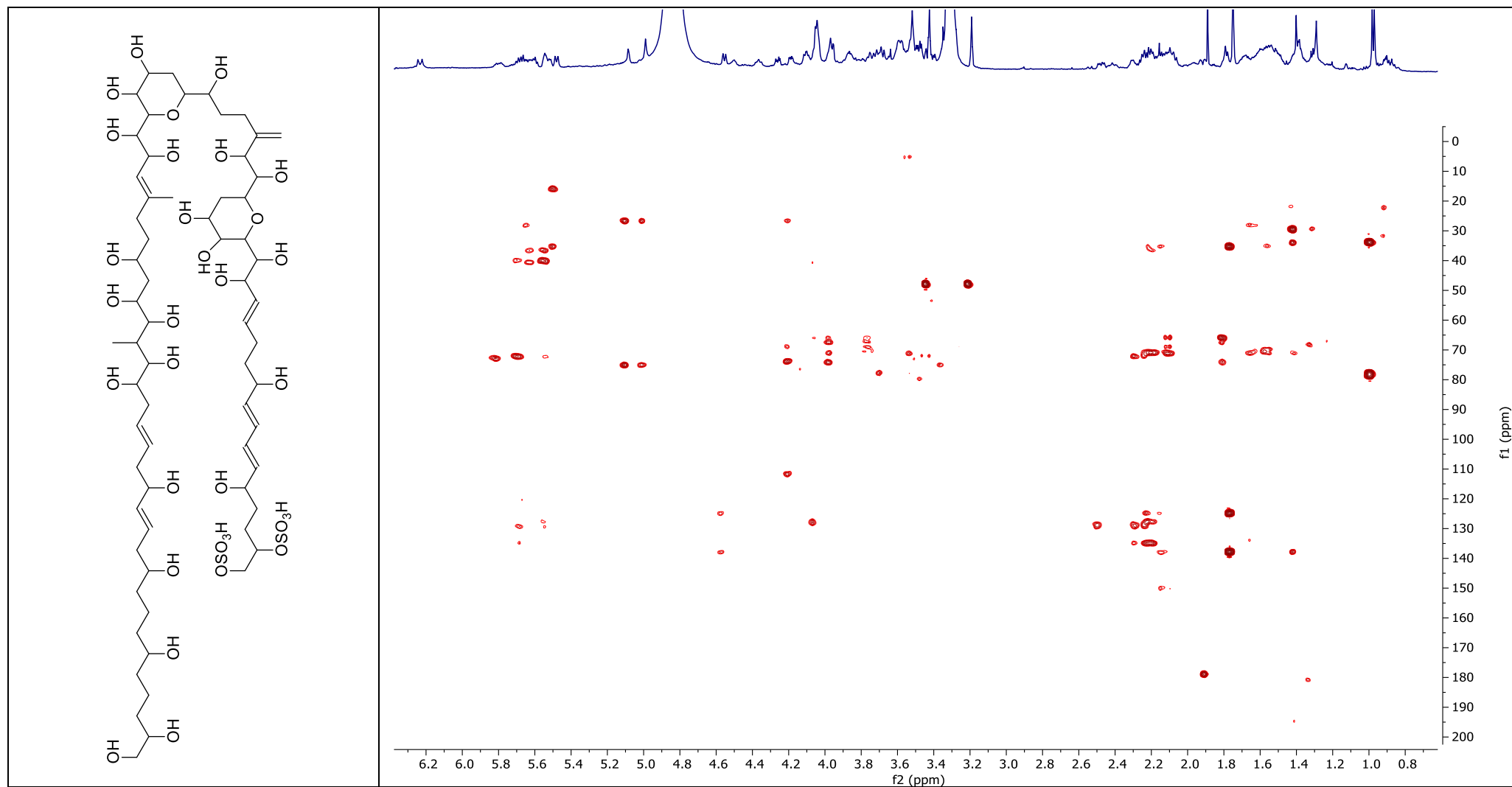


Figure S23. HRESIMS spectrum for amphidinol 25

Elemental Composition Report

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -50.0, max = 75.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Odd and Even Electron Ions
1111 formula(e) evaluated with 31 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 5-80 H: 0-150 O: 0-40 Na: 0-1 S: 0-2
18_04_11 Sulfatado NEG 161 (0.694) Cm (159:177)
1: TOF MS ES-

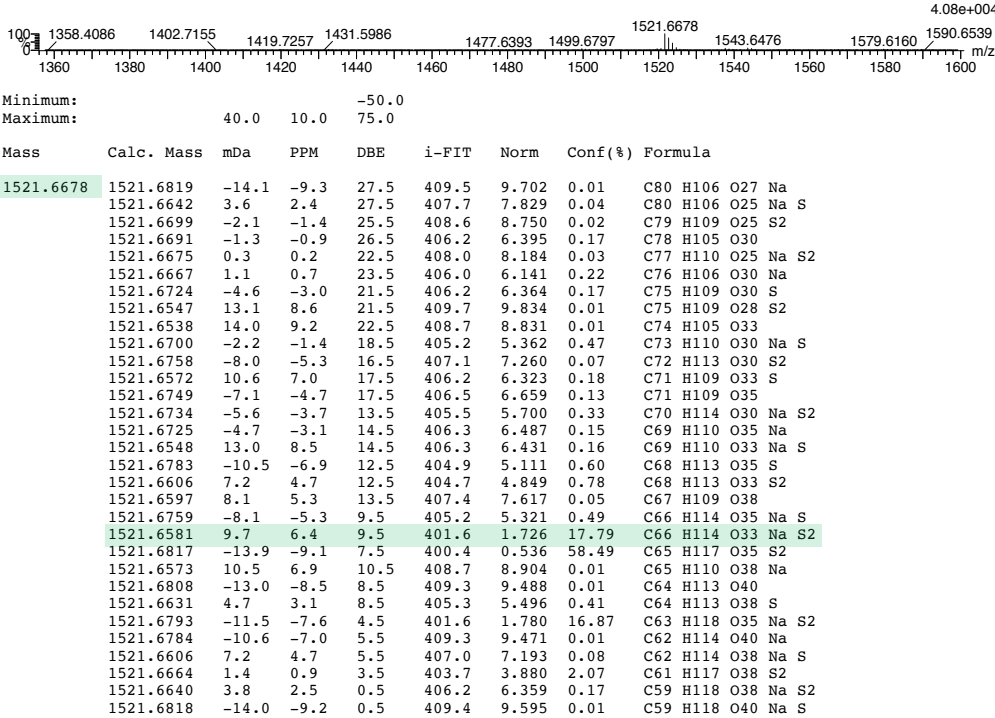


Figure S24. Main MS/MS fragments observed for amphidinol 25.

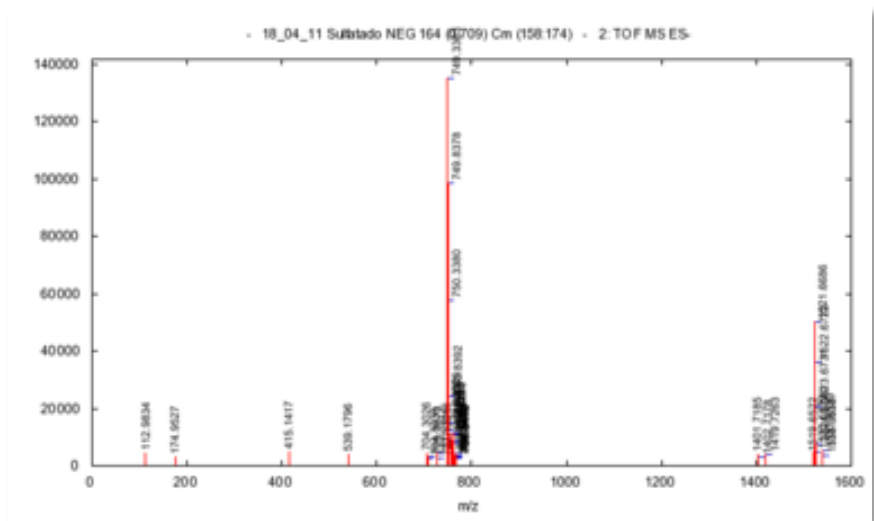


Figure S25. ^1H NMR spectrum (600 MHz, CD_3OD) for amphidinol 26.

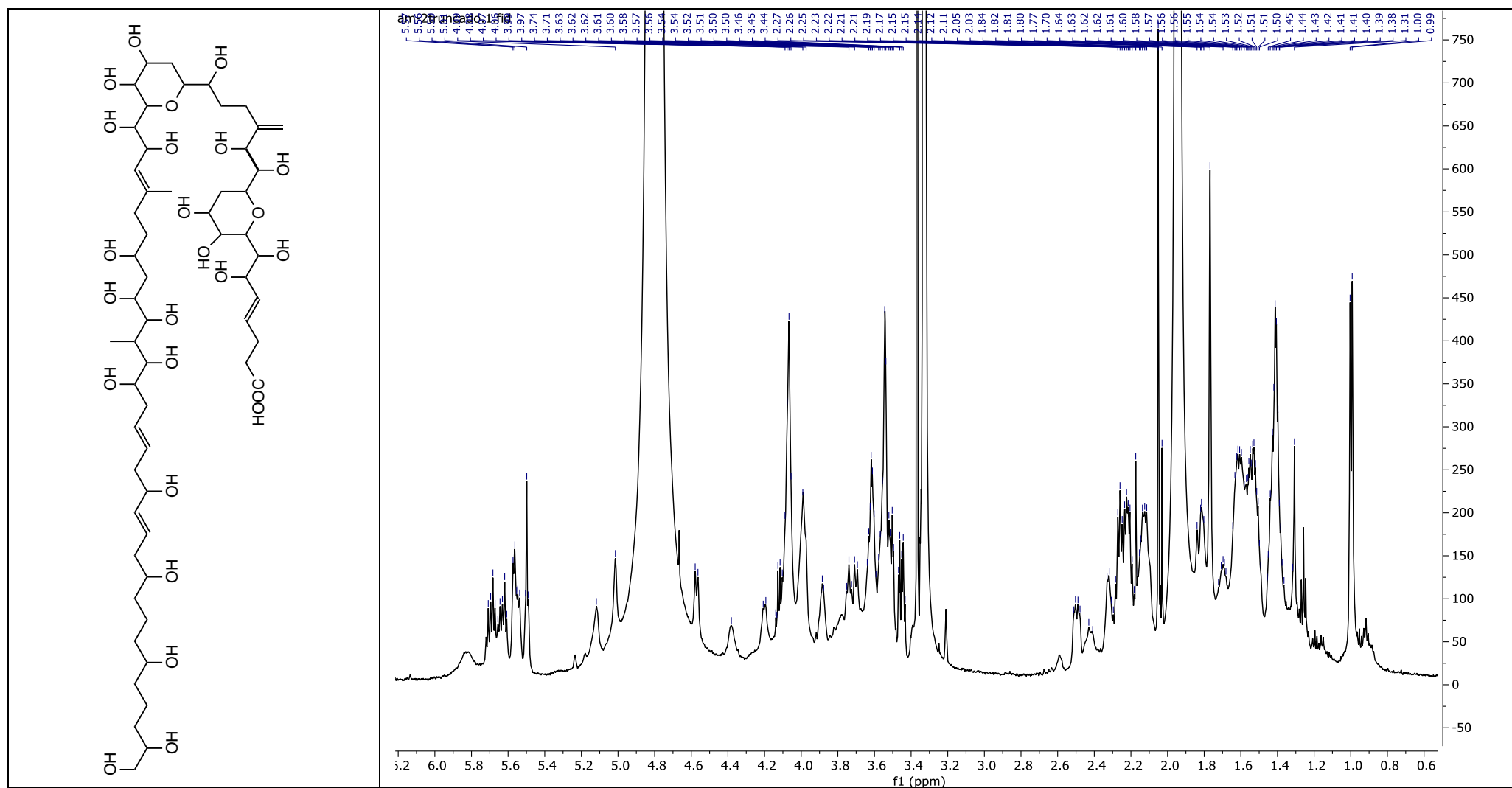


Figure S26. COSY spectrum (600 MHz, CD₃OD) for amphidinol 26.

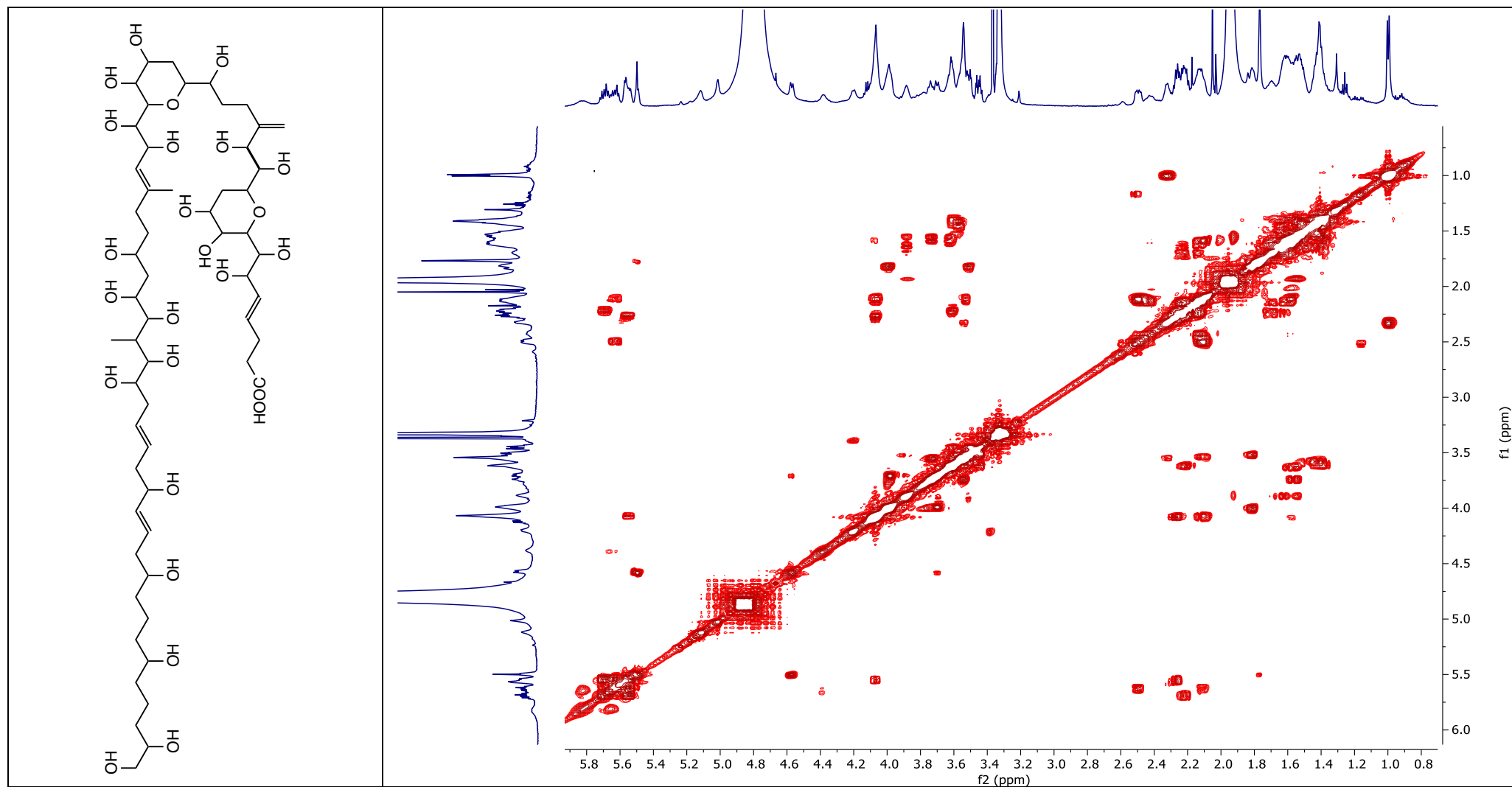


Figure S27. HSQC_{ed} spectrum (600 MHz, CD₃OD) for amphidinol 26.

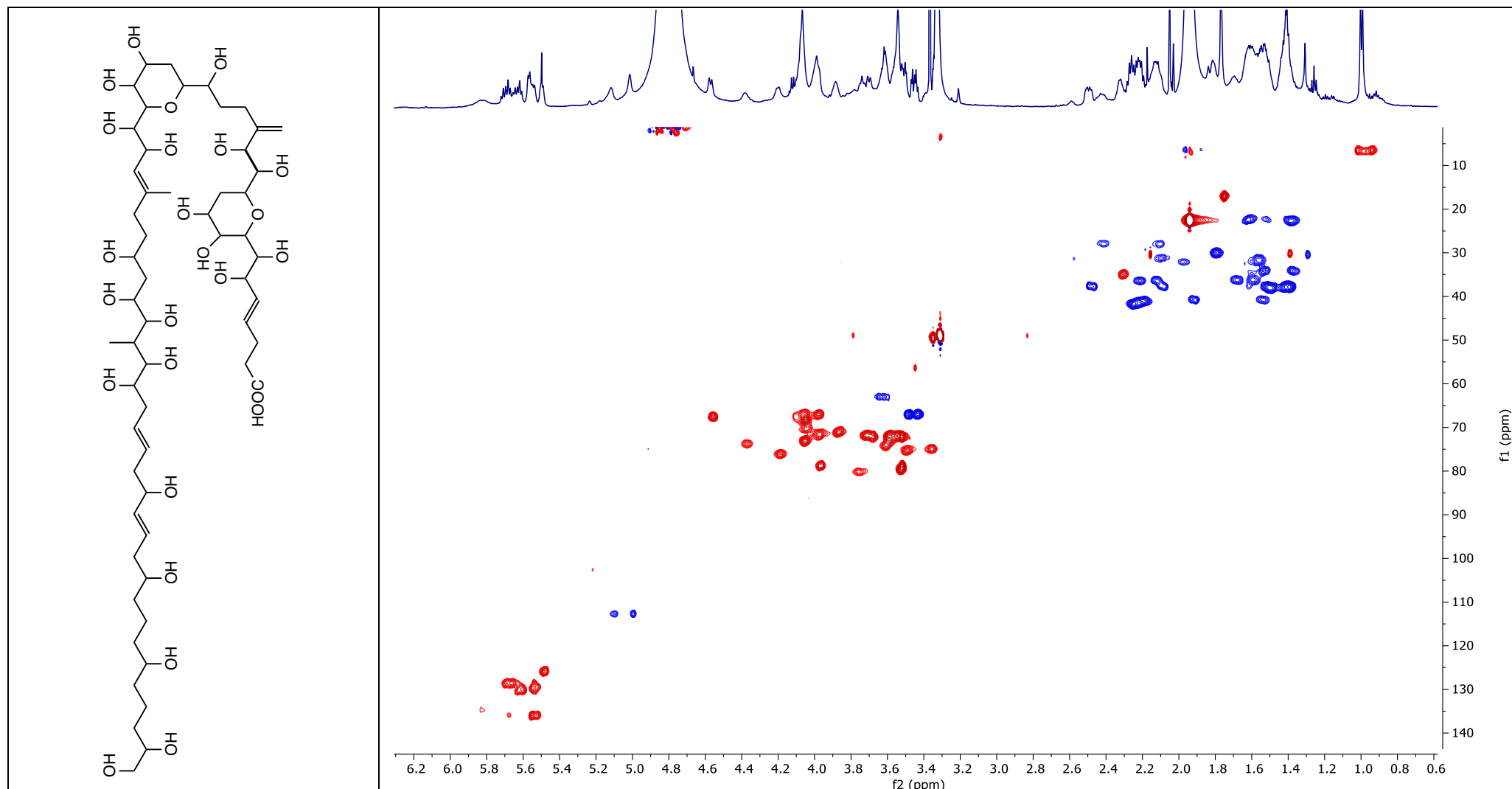


Figure S28. HSQC-TOCSY spectrum (600 MHz, CD₃OD) for amphidinol 26.

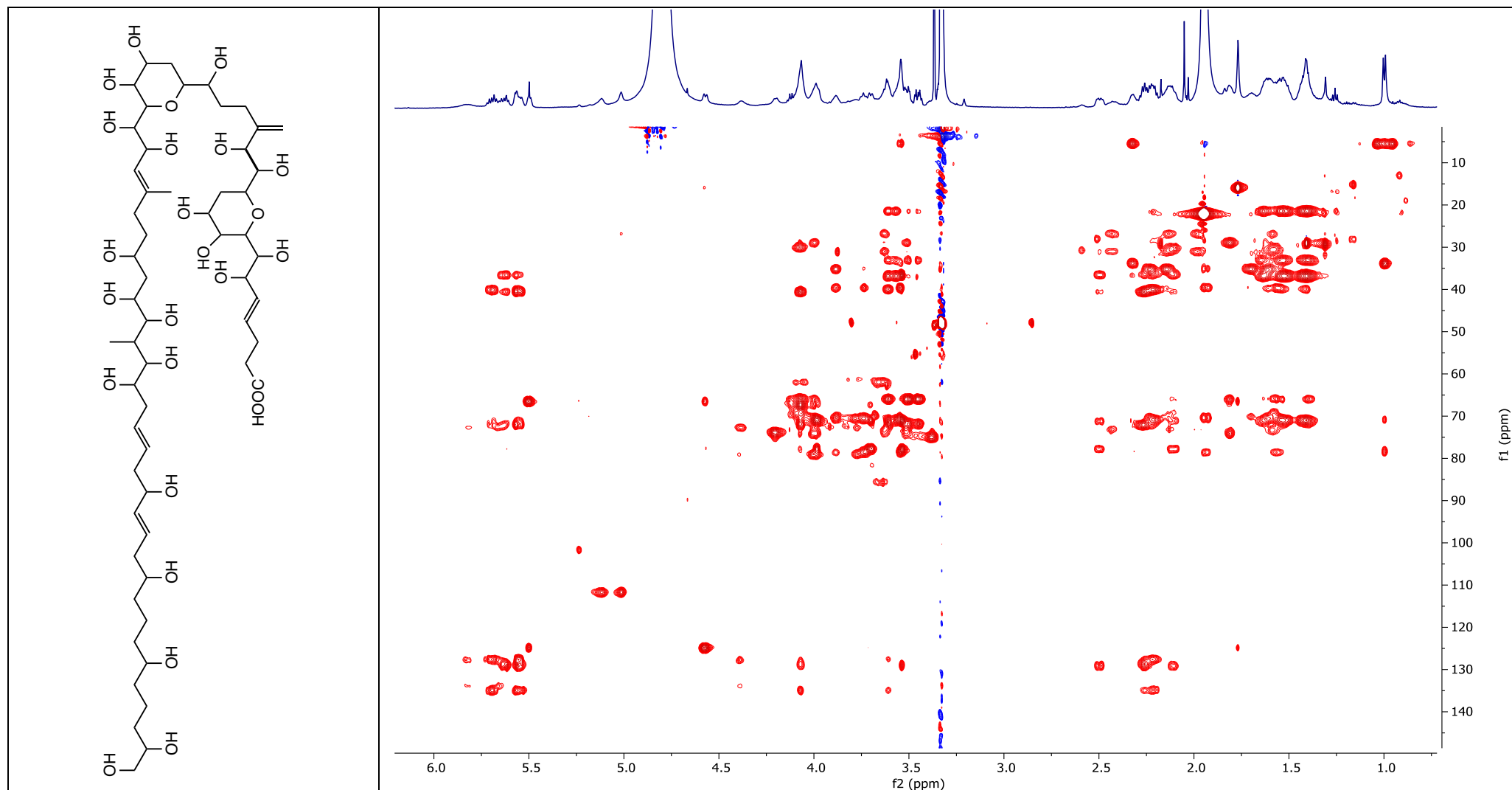


Figure S29. HMBC spectrum (600 MHz, CD₃OD) for amphidinol 26.

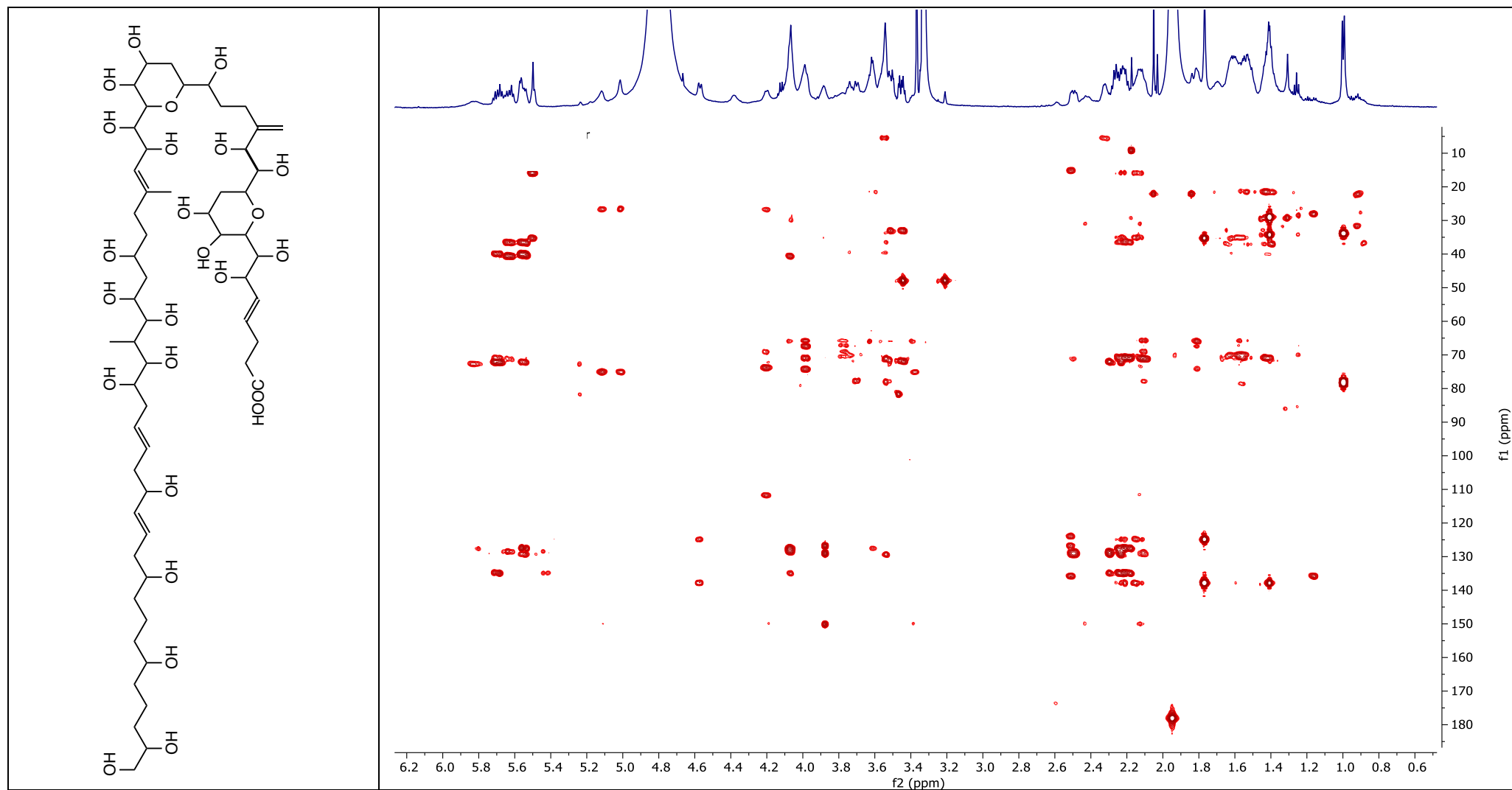


Figure S30. H2BC spectrum (600 MHz, CD₃OD) for amphidinol 26.

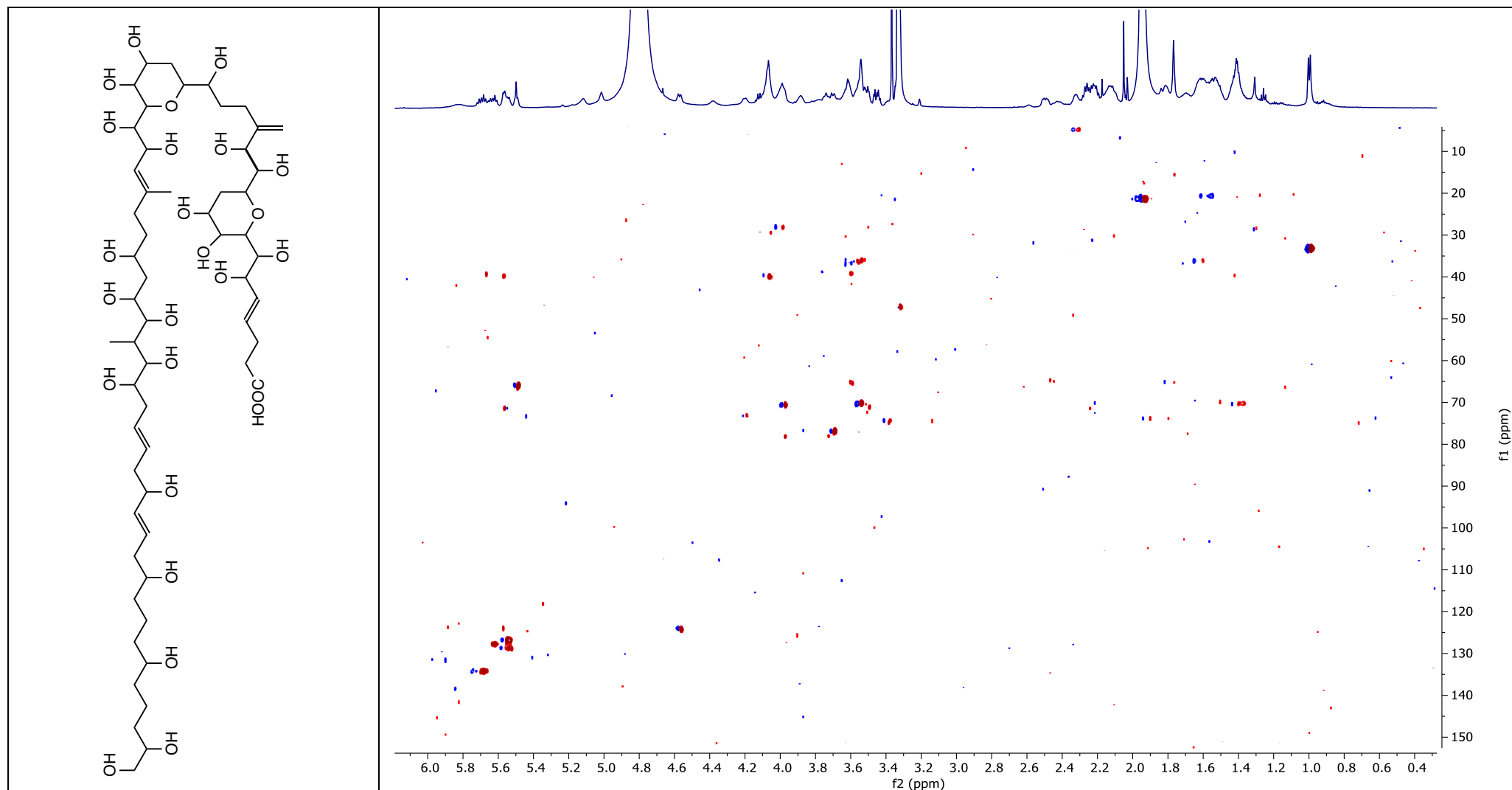


Figure S31. T-ROESY spectrum (600 MHz, CD₃OD) for amphidinol 26.

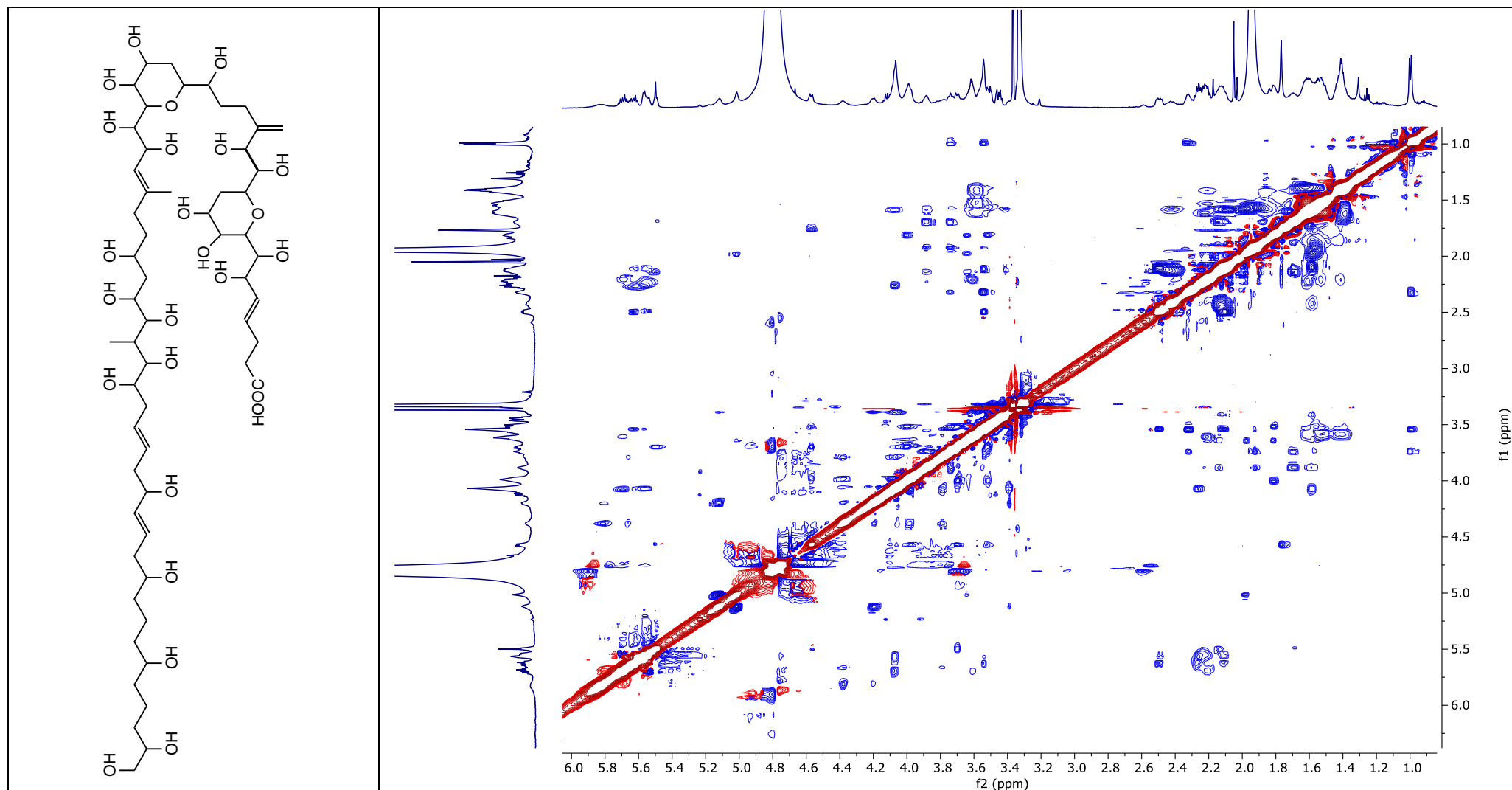


Figure S32. HRESIMS spectrum for amphidinol 26.

Elemental Composition Report

Page 1

Multiple Mass Analysis: 3 mass(es) processed

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 120.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

325 formula(e) evaluated with 5 results within limits (all results (up to 1000) for each mass)

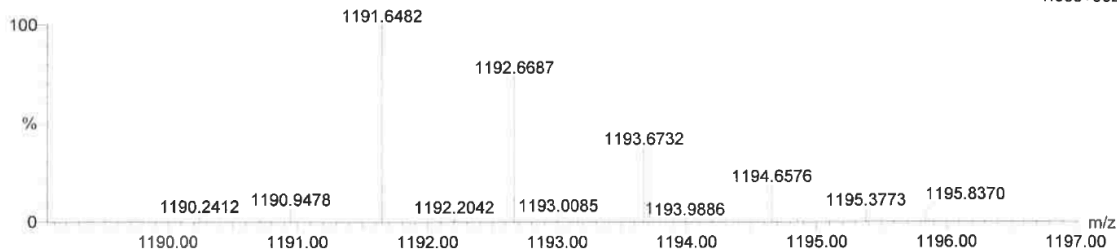
Elements Used:

C: 0-60 H: 0-110 O: 0-25 Na: 0-2

David

ESI (17-159) Adrian (ACO35CI8MAC) 51 (1.778)

2: TOF MS ES+
1.68e+002



Minimum: 20.00

Maximum: 100.00

5.0 5.0 -1.5

120.0

Mass	RA	Calc. Mass	mDa	PPM	DBE	i-FIT	i-FIT (Norm)	Formula
1191.6482	100.00	1191.6526	-4.4	-3.7	10.5	26.5	0.5	C59 H99 O24
		1191.6502	-2.0	-1.7	7.5	27.2	1.3	C57 H100 O24 Na
		1191.6478	0.4	0.3	4.5	28.1	2.2	C55 H101 O24 Na2
1192.6687	73.82	---	---	---	---	---	---	---
1193.6732	37.73	1193.6683	4.9	4.1	9.5	33.5	0.4	C59 H101 O24
		1193.6787	-5.5	-4.6	7.5	34.4	1.2	C59 H103 O21 Na2

Figure S33. Amphidinol 26 conversion from aldehyde to carboxylic acid at C-54 observed by ESI-HRMS.

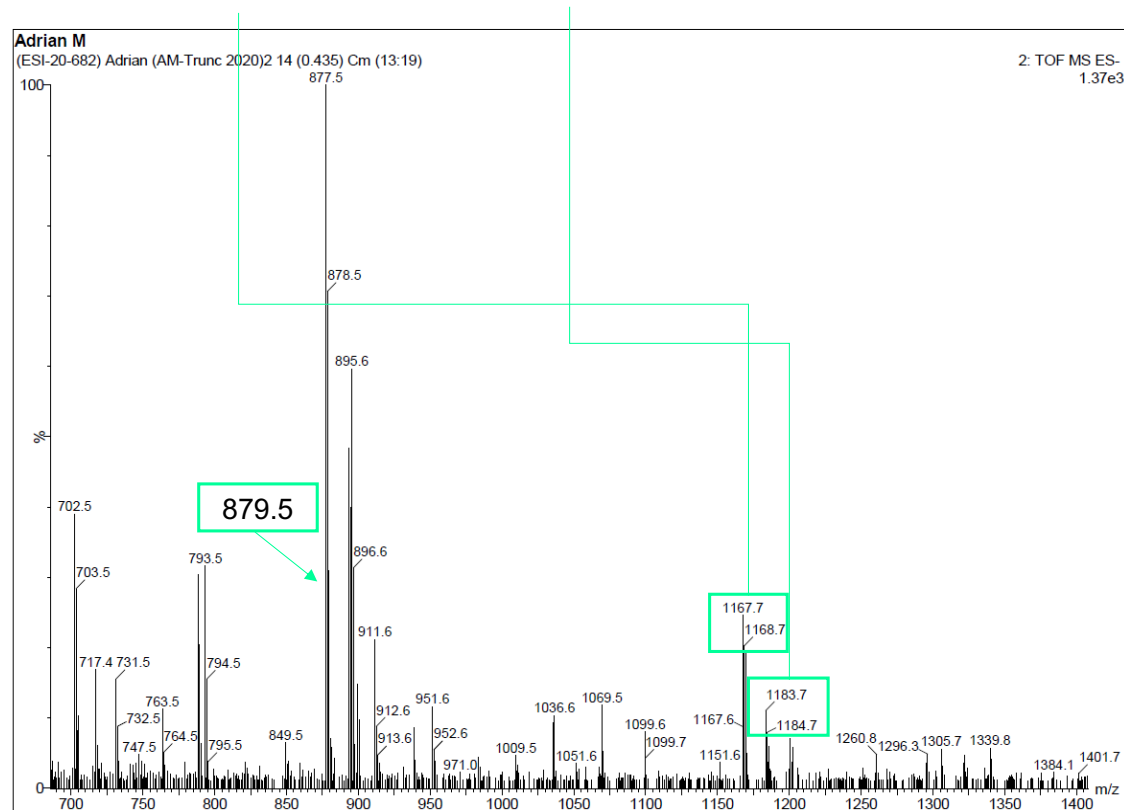
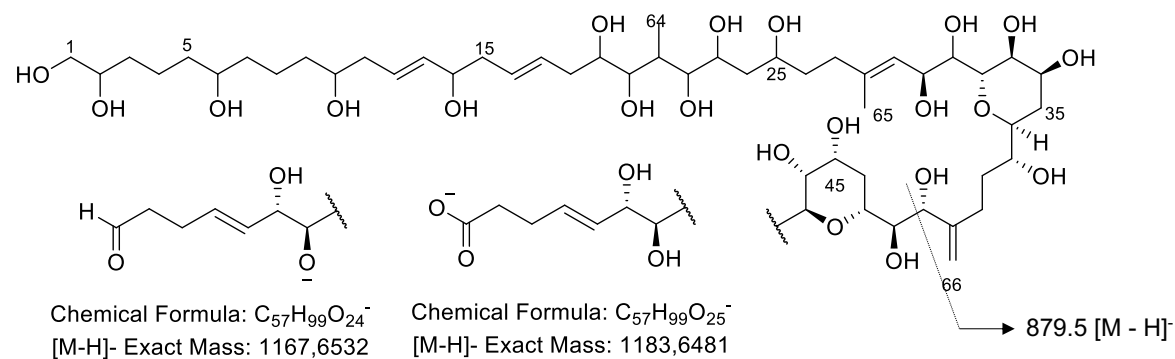


Figure S34. Amphidinol 26 single mass composition analysis for aldehyde and carboxylic states.

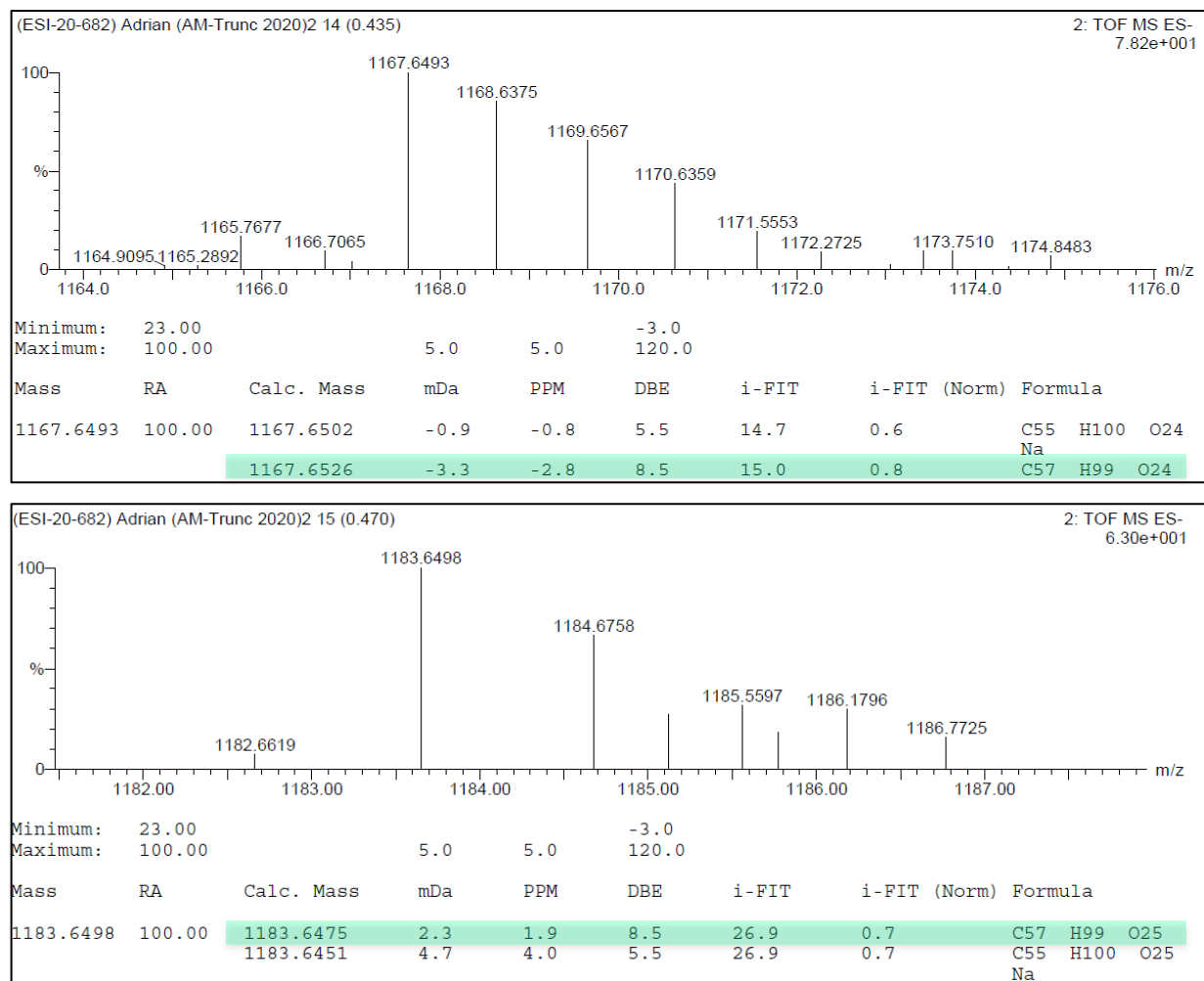


Figure S35. Main MS/MS fragments observed for amphidinol 26 as aldehyde.

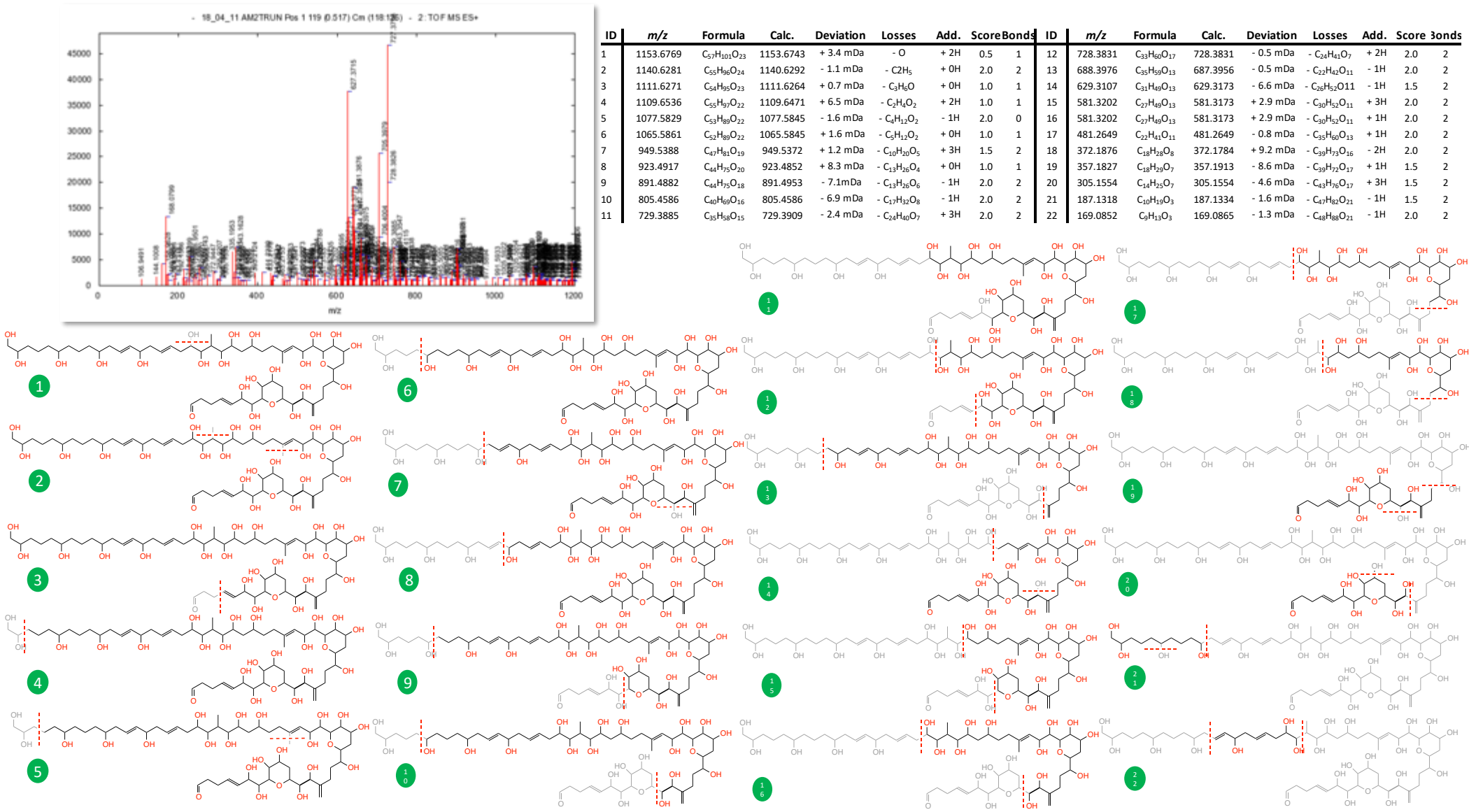


Figure S36. ^1H -NMR and HSQC_{ed} spectra (600 MHz, CD_3OD) for luteophanol D.

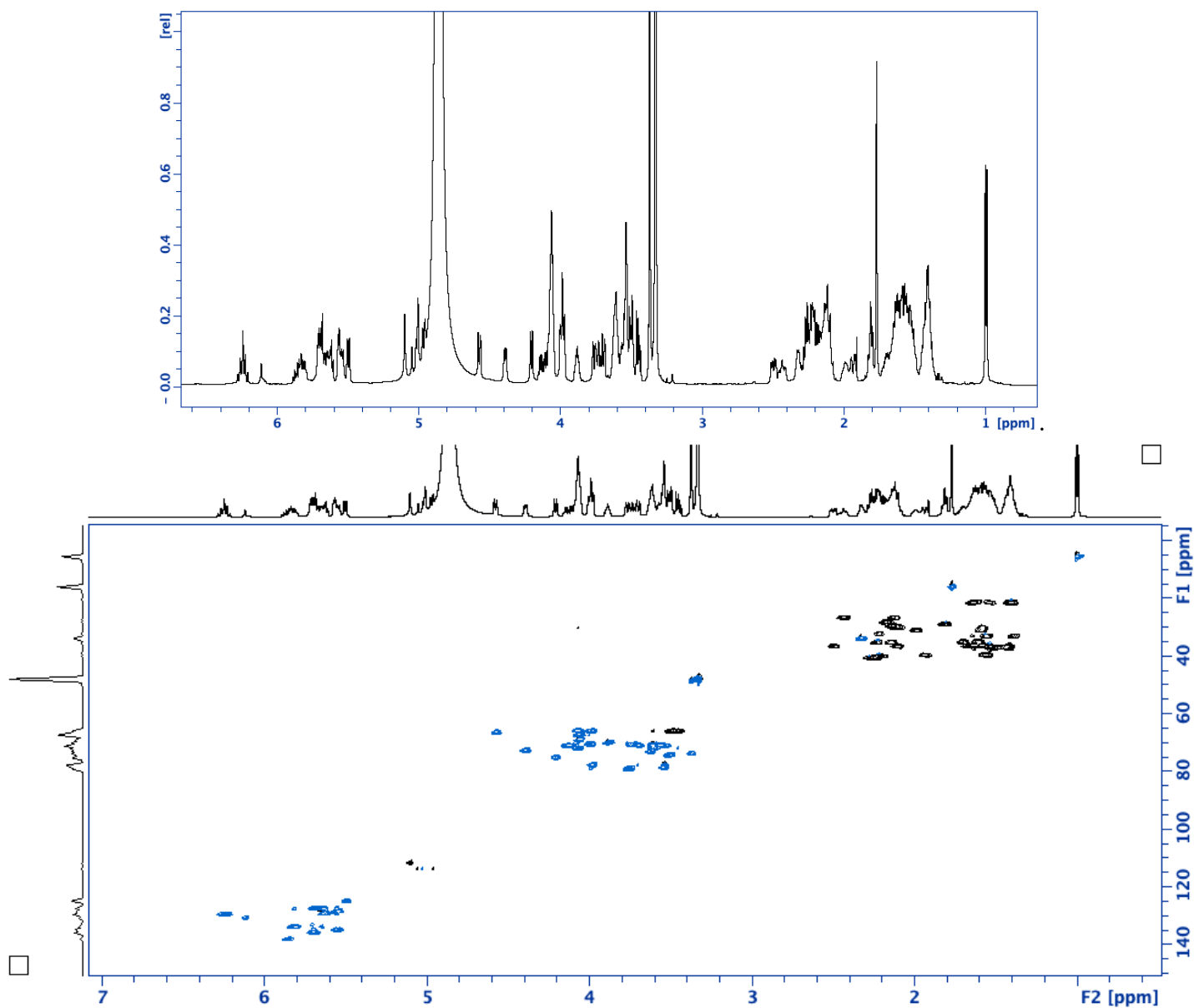
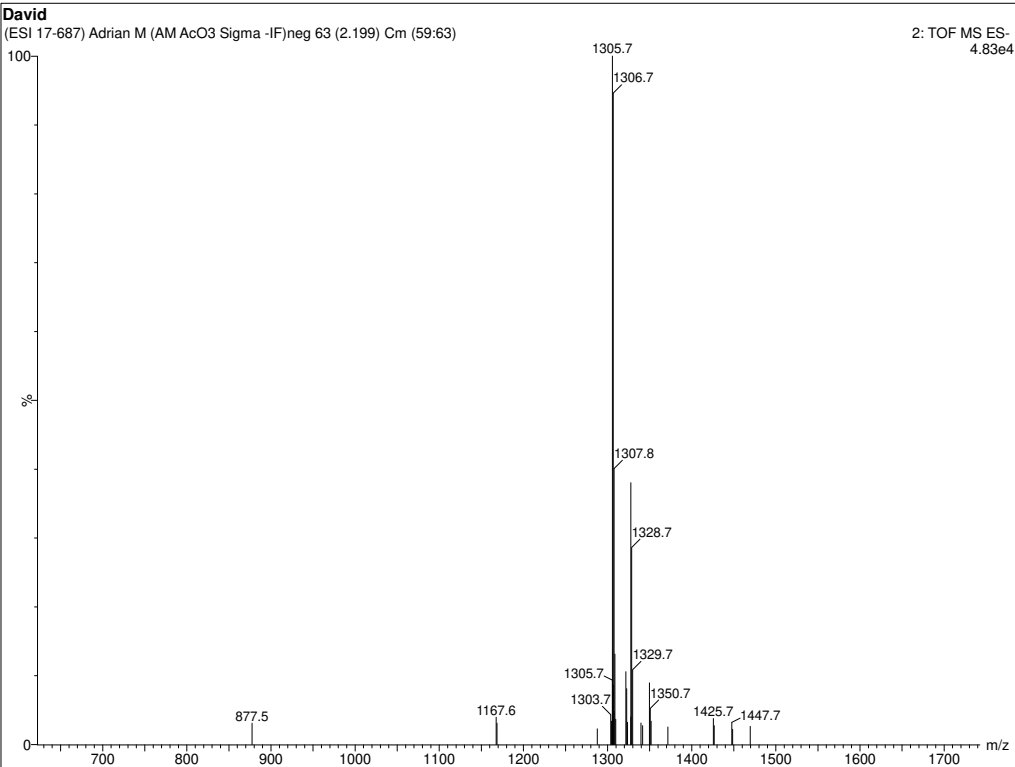


Figure S37. HRESIMS spectrum for luteophanol D.



Elemental Composition Report

Page 1

Multiple Mass Analysis: 3 mass(es) processed
Tolerance = 5.0 PPM / DBE: min = -1.5, max = 100.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
906 formula(e) evaluated with 9 results within limits (up to 50 best isotopic matches for each mass)
Elements Used:
C: 0-80 H: 0-120 O: 0-30 Na: 0-1
David
(ESI 17-687) Adrian M (AM AcO3 Sigma -IF)neg 94 (3.279)

2: TOF MS ES-
5.24e+003

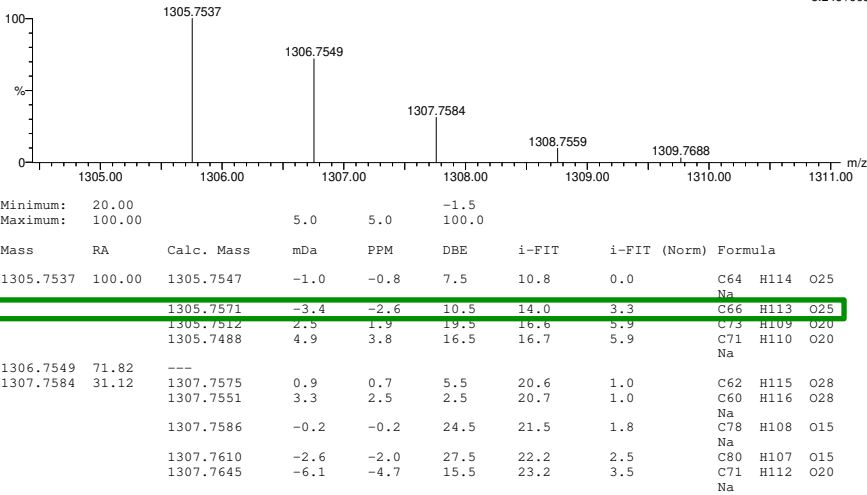


Figure S38. ^1H -NMR and HSQC_{ed} spectra (600 MHz, CD_3OD) for amphidinol 20B.

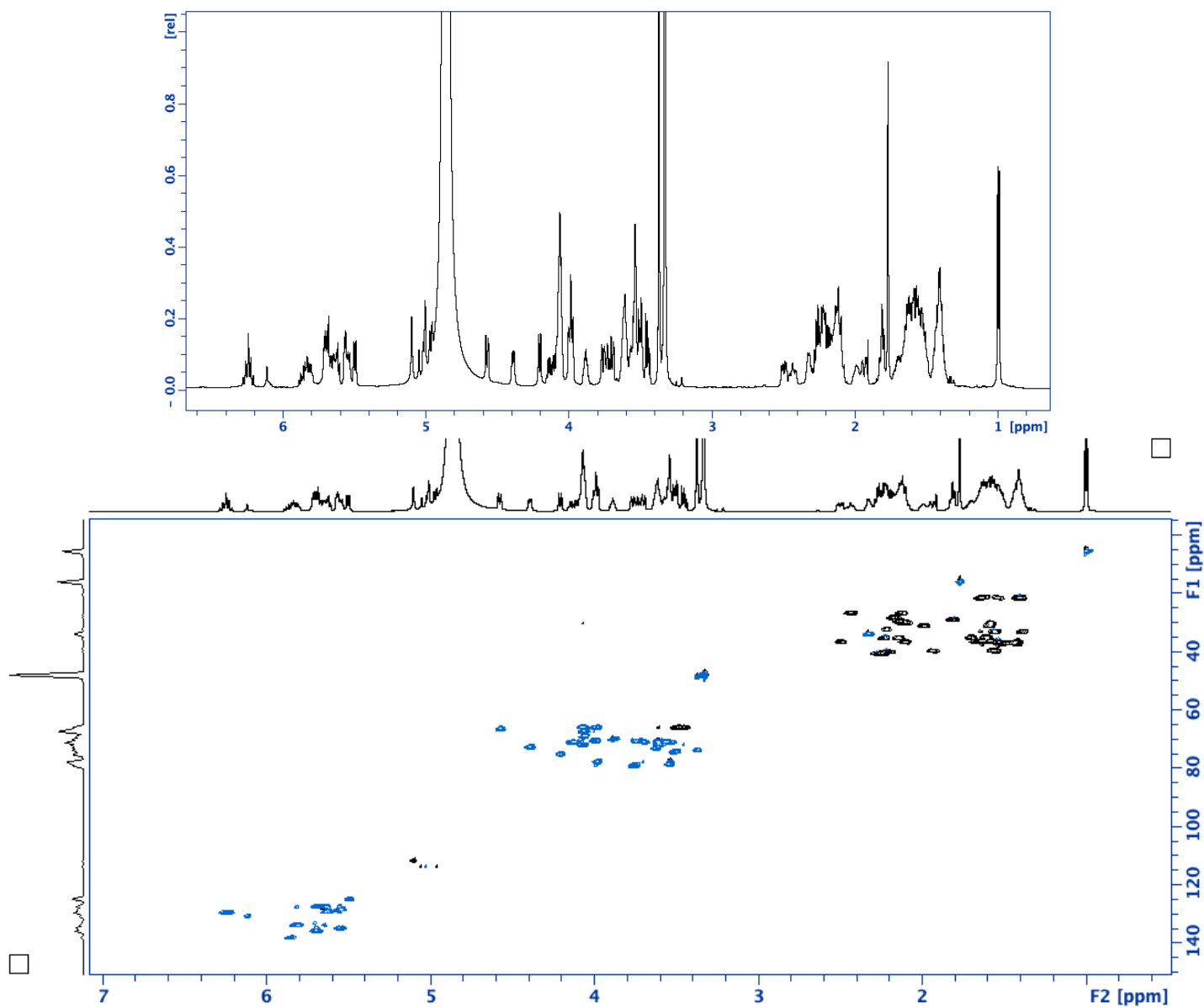
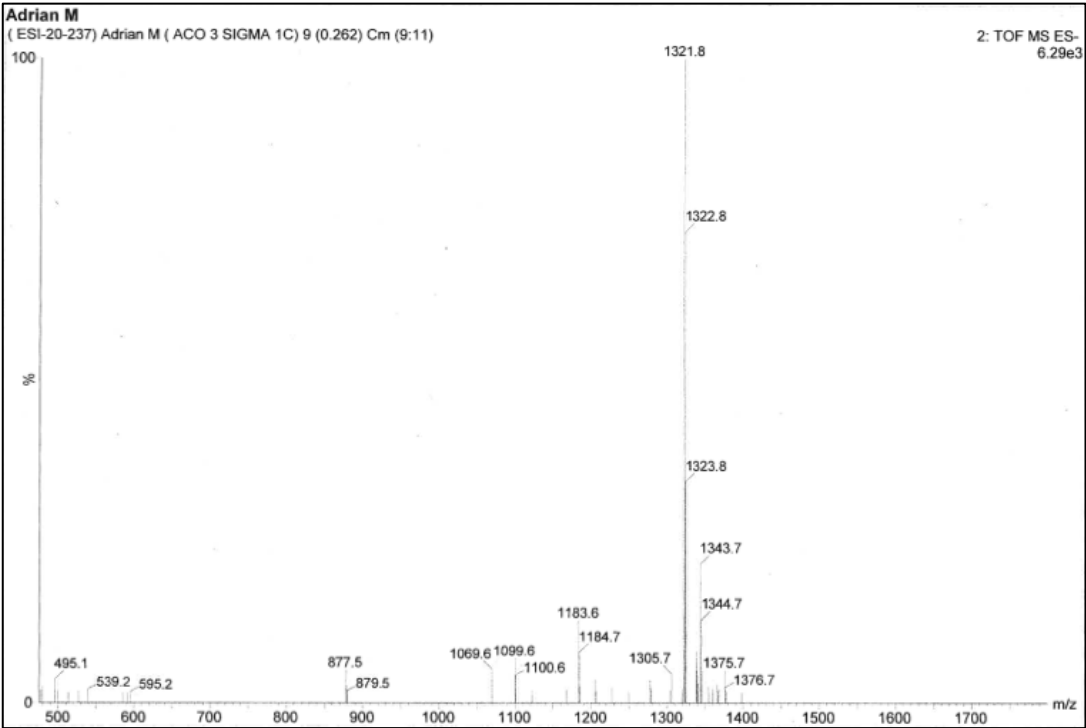


Figure S39. HRESIMS spectrum for amphidinol 20B.



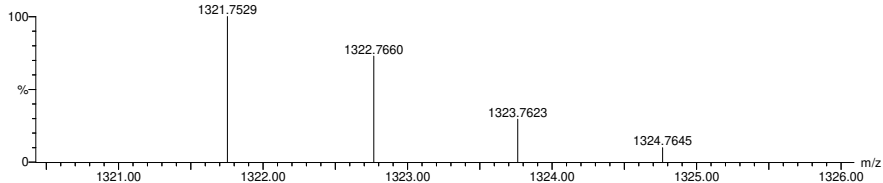
Elemental Composition Report

Page 1

Multiple Mass Analysis: 3 mass(es) processed
Tolerance = 5.0 PPM / DBE: min = -1.5, max = 100.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
847 formula(e) evaluated with 8 results within limits (up to 50 best isotopic matches for each mass)
Elements Used:
C: 0-80 H: 0-120 O: 0-30 Na: 0-1
Adrian M
(ESI-20-237) Adrian M (ACO 3 SIGMA 1C) 34 (1.462)

1: TOF MS ES-
9.11e+002



Minimum:	20.00				-1.5				
Maximum:	100.00		5.0	5.0	100.0				
Mass	RA	Calc. Mass	mDa	PPM	DBE	i-FIT	i-FIT (Norm)	Formula	
1321.7529	100.00	1321.7496	3.3	2.5	7.5	13.1	0.7	C64 H114 O26	
		1321.7520	0.9	0.7	10.5	13.2	0.8	C66 H113 O26	
		1321.7590	-6.1	-4.6	20.5	15.8	3.4	C75 H110 O18	
								Na	
1322.7660	72.80	---							
1323.7623	29.39	1323.7653	-3.0	-2.3	6.5	22.9	1.5	C64 H116 O26	
		1323.7618	0.5	0.4	18.5	23.0	1.5	Na	
		1323.7594	2.9	2.2	15.5	23.0	1.6	C73 H111 O21	
								C71 H112 O21	
								Na	
		1323.7677	-5.4	-4.1	9.5	23.1	1.6	C66 H115 O26	
		1323.7559	6.4	4.8	27.5	23.3	1.9	C80 H107 O16	